SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
:	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
3863	34231	A	3903	174	1599	VLHAVGNQVQGCPGPEVHVCG
		İ				WRPGCVRLLQLHVVGRADGPE
						CDLCAVWPGGGDV*PAVPGAV
						PHVTQCEPQGHPYGEGTGAGA
						YCAALCGPPGARGHCGQEARQ
		ŀ				PQVRTCQGQERRRDCQDLL*E
						AGGQEAPGAERPSAG*ASRGTP
						RAAATHPPRTASPGEGQHV*VP
						AHVGGARPGAWQRHPGLHQY
						HRPQHLEPPAQPDEPHQDP*HL
		1				PGQRPAPAVEAEPGQQAPS\KP
						CPPPEPSAPQDGVPAENGLPQG
						DP/GAIAPRAQAPDSPCGRCTSP
						GQQ*YWLGPGAPQRGSSPEWD
						RP*ATQDGRPRPRPTPAAAICDP
						G*PREPRGGAPQWAGWGGRRR
		ļ				**RLRNPQ*PGQP\SGSSGRGPG
						PRRPSVASSVSE/RVLRGERALS
						PSPEAPLRASGQRANPTTAPAA
						ECPPYNPRDLCWTPGWLPMGP
		ļ				ESGKRRSRCVEEDAGPALHRQ
		ļ				GGTDGET**TGRGGNRPGPYGR
3864	34232	Α	3904	331	1120	HKDRFWQLQNDSCFLHSPGER
						QWLGGPRSDTFGPQVLFGHVGI
]						CSQRA/HPAGPGHRGLPEGR*PP
						HRSQRHPPRSRKPYLA*PPDMC
						VATDRRTQTPRDFPPLGR*KPH
						GTLRSAACPAGRVSPSPRPRGL
ļ						PAPPPKSHLCG\PGVRGR*QLLP
						PHPGSPKGERGWTASPGAARG
		ł				GPGPAPAPRP\RASWSQPSVTFP
						LPLAGLA/GHPGSRTEPAWKAG
Ì						GAAARPGPELPRDLLQAGSTDT
						ASGEQLAAGPWTGKEISGRARP
	-]		RL
3865	34233	Α	3905	2	415	YTILTEK*KLSKLST*W\VHQDQ
						LQKREELSMEILNK*DQDSEAY
						PQRTVTGEETWLYQYDPPPLPR
1						SLPPPQTHTAPVGA*S/DWGG*E
						LPPGLNGDKLAHHSPTPFLSFSG
						LLFVDWL*SQLLSLFGLFTQGVI
						RIFI

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
}				sequence		
3866	34234	Α	3906	1	4527	MGFCCRECRPRLLKGRPCIQHA
		1				GPVAAFKVATPYSLYVCPEGQ
l						NVTLTCRLLGPVDKGHDVTFY
						KTWYRSSRGEVQTCSERRPIRN
]		Ì	LTFQDLHLHHGGHQAANTSHD
		1				LAQRHGLESASDHHGNFSITMR
						NLTLLDSGLYCCLVVEIRHHHS
		1		,		EHRVHGAMELQVQTGKDAPSN
						CVVYPSSSQDSESNHGNNFRIH
	•					VSNGLLMRGPRPLDRERNSSHV
			:			LIVEAYNHDLGPMRSSVRMRK
						LRQSTALAQHWTGTALDR
3867	34235	Α	3907	1	2180	MALTFPCRKFEWYGRRQPEVR
		1				YSVPASHQLKATDADEGEFGR
		1				VWYRILHGNHGNNFRIHVSNG
					,	LLMRGPRPLDRERNSSHVLIVE
						AYNHDLGPMRSSVRMRKLRQS
						TAL\DSTGQAQHWTESRSGSPG
						SPVAPTCSART*QTSAS\VHLCL
						SGKSHHAWPP*TPFKLYYVH\E
		ŀ				YSAHIHKENLVLVIVYVEDIND
						EAPVFTQQQYSRLGLRETAGIG
						TSVIVVQATDRDSGDGGLVNY
						RILSGAEGKFEIDESTGLIITVNY
						LDYETKTSYMMNVSATDQAPP
						FNQGFCSVYITLLNELDEAVQF
						SNASYEAAILENLALGTEIVRV
	1					QAYSIDNLNQITYRFDAYTSTQ
						AKALFKIDAITRILGTQMDTKM
						NKTLLSPQRVLRLEVEMELIQD
			•			ANQSATRRCAENYNRGVVEPL
						RAQQSYLAGEAGRLHGRGGFP
						VECEREEGIQQTECPGEVMPDR
						GSDMEGVITVQGLVDREKGDF
						YTLTVVADDGGPKVDSTVVSG
						TRVYITVLDENDNSPRFDFTSDS
						AVSIPEDCPVGQRVATVKAWD
						PDAGSNGQVVFSLASGNIAGAF
						EIVTTNDSIGEVFVARPLDREEL
						DHYILQVVASDRGTPPRKKDHI
						LQVTILDINDNPPVIESPFGYNV
						SVNENVGGGTAVVQVRATDRD
						IGINSVLSYYITEGNKDMTFRM
						DRISGEIATRPAPPDRERQSFYH

SEQ ID	SEQ ID NO:		SEQ ID NO:		1	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
ļ				sequence	or peptial sequence	determine, v possible intercounter insertion)
	<u> </u>	<u> </u>			ļ <u>.</u>	
3868	34236	Α	3908	603	1395	RGRPRGSPIFPTRAPREKKR*ÉE
						VGGHKREQTG*GGGERRKPPN
					1	PQHEPKERGWCSRVPEEPQ/RK
						RRSARARPKKL*REKRRRGRPK
						RCLW*TGRHPSHHPRTHQC*F*
						WRK/REEGKERKKEQPAHAGQ
				ļ		KRRKAARHRRRRRRRRERTDEK
			•			NRTWTRRRREKAGQDEKREGE
				•		HGQKRSQQGRESRRDGRARTR
						KERRQKRENDNRARRRQQAER
		1				EKTKSVKRRQTTQQAEEVRQA
		l				RENEAREPQQRRQHSRRRKEKE
		<u> </u>				EMRAPRSKQ
3869	34237	A	3909	1	548	
3870 3871	34238	A	3910	1	1803	
3871	34239 34240	A	3911 3912	1	279 506	MCVCDOCNI CTECECKINGCEV
38/2	34240	A	3912	1	300	MCYSRQSNLGTFGEGKIKGSEV
						IDECPRSSRYQDLQELQNKTKL
		l				TVLEGDILDESCLKRACQDMSV
						IIHTTSIIDIIGVTHRESIMNINVK
				1		RTQLLLEACVQATVPVFIYTSTP
		İ				EVAGPNSYKEIIQNSHEEEPLEN
	:					T\WCSPS/PYKKA/LARSGI*ATL
2972	24241	_	2012	12	(21	QLGGSQEECT
3873	34241	Α	3913	3	621	AGQQTVEIDLRHRIQLP\DLENQ
İ		·				RNFNELSRIVLEVRERVRQEQQ
						EGGHEAGEGRGRQGPRESQPSP
						AQPRAEAPSKGPDGTPGEDGGE
						PGDAVAAAEQPAQCGQGQPFV
]						LPVGVSSRNEDYPRTCRMWNS
						TFQTYKKEVCLPRHSMHPGPW
	ŀ			•		AICCECQTRFGGRLPVSRVEAA
						LPYWVPLSLRPRKQHPCWMHA
3874	34242	A	3914	1	420	AGTTAGGSAVMS
38/4	34242	A	3914	1	430	RHRIQLPDLENQRNFNELSRIVL
						EVRERVRQEQQEGGHEAGEGR
						GRQGPRESQPSPAQPRAEAPSK
•						GPDGTPGEDGGEPGN\AVAAAE
						QPAQCGQGQPFVLPVGVSSRNE
						DYPRTCRMW*GCGGYWGLKV
3875	34243	Α	3915	2	1175	GQHGLQRGPQPHT
3876	34244		3915	1		HLRIHTQESSYVCDECGKALTS
3070	27277	^	3710	1		KRNLHQHQRIHTGEKPYECSKY
]			,	\G*PFGLLPQLGHLEHVYSGEKP
						VLDICRFGLPEFFTPFYW
L	<u> </u>	L				VEDICALGUETETTA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
3877	34245	A	3917	<u> </u>	1396	MRPQLAGRDHHRGAATLLLER
	13 ,2 .3	` `		•	1.070	PGRLVTHFRQRRGAVRYGGGK
	1					STTQHLSQRRLSGPNDHTKGLV
						WLLEHILQRLVSFVKLQATRTF
						TRTYITYAWFLPWGFSGVLCGT
		1				PVDTCWALKHQRIHTGEKPFEC
		İ				SECGKAFNGNSSLIRHQRIHTGE
						RPYQCEECGRAFNDNANLIRHQ
						RIHSGDRPYYCTECGNSFTSSSE
		1				FVIHQRIHTGEKPYECNECGKA
						FVGNSPLLRHQKIHTGEKPYEC
						NECGKSFGRTSHLSQHQRIHTG
						EKPYSCKVCGQAFNFHTKLTR
						HQRIHSEEKPF*L/CVDCGKAFS
						AQEQLKRHLRIHTQESSYVCDE
						CGKALTSKRNLHQHQRIHTGE
						KPYECSKYEKAFGTSSQLGHLE
						HVYSGEKPVLDICRFGLPEFFTP
		1				FYWKEEKKCGRKMRNEVVHK
		1				VSFFLVVPIALSSLLKKKWKML
	j					KKEKAQDPTEYGNLEDDNSQQ
3878	34246	A	3918	1	547	MDSQRPRER/QRERERQSERQR
3676	34240	^	3710	*	547	HTQRMHREAETEDERDWKGH
						DTKTRRQRQRKRAEEGQCREH
						DRERRRD\RTGERREKQRKSTQ
						QSRKPSEEPHREKTQIKRERGPE
						QGELERGQCTERNRKA/GTPEC
ŀ						*TDPHIWTPHPARSAPAHPPDH
						TAAKYRPPYRSHHSGITHQHPR
						AASTLKLWPKP
3879	34247	Α	3919	1	399	THE TELLE WITH
3880	34248	Α	3920	3	872	KSKLKSEQDGISKTHKLLRRTC
						SSTVKTDDVCVTKSHRTFGRSL
	İ					SSDPRAEQAMTAIKSHKLLNRP
						CPAAVKSEECLTLKSHRLLTRS
						WSGDPRCEHNTNLKPHKLLSRS
						YSSNLRMEELYGLKNHKLLSKS
						YSSAPKSSKT*/VFSKEP*RRRG
						RKALSLPQGLFGYP*HHLHPSSS
						QLAPNGAKCIPVRDRGFLVQTI
	:					EFAEQRIPVLNEYCEVCDEPHV
						FQNGPMLRRGRDVCEWAKKY
						ANSVVRKKFCRLSIARRSRYRA
						DMDLLRMSNFILTIIYKQKLNL
3881	34249	Α	3921	3	218	CCRSHQGAGEGGHLSVQLLWQ
						YRWMWCSWGPV\FQFHTDLEL
						VAWRCVGLDPGCQQLDIGMQT
						LIGDHICVF
3882	34250	Α	3922	1	1055	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
3883	34251	A	3923	3	962	RSMRQKVNKDIQDLISALPQAD LIDIYRTLHPKSTEYTFFSAPHR TYFKIDHIVGSKALLSKCKRTEI TTNCLSDHSAIKLELRIKKLTQN RTTTWKLNNLLLNDYWVHNE MKAEIKMLFETNENK/DETYQN LWDTFKA/PSIILNGQKLEAFPL KTGTREGYLLLPLLFNIVLEVL AMAIRQEKE/IKGFQLGKEEVK LSLFADDMIVYLEDPIISAANLL RLISNFSKVSGYKINVQKSQTFL YTNNRQTESQIMSELPFTIATKR IKYLGIQLTRDVKDLLKENYKP LFNEIKEDTNKWKNIPCSWIGRI KIVKMAILPK
3884	34252	A	3924		1452	MGDFNTPLSTLDRSMRQKVNK DTQELNSALHQADLIDIYRTLH PKSKEYTFFSALHHTYSKIDRT VGSKALLSKCKRTEIITNSLSDH RAIKLELRIKKLTQNRSTTWKL NNLLLNDYWVHNEMKAEIKM FFETNENKDTTYQNLWDTFKA VCRGKFIALNAHNRKQERSKID TLTSQLKELEKQEQTQSKASRR QEITKIRAELMEIETQKTLQKSN ESRSWFFERINKIDRPLARLIKK KREKNQIDVIKNDKGDITTDPT EIQTTIREYYKHLYANKLENLE EMDKFLDTYTLPRLNQEEVESL NRPITGSEIVAIINSLPTKKSPGP DGFTVEFY/QEGN*AGEGNKGY SIRKRRSQIVPVWR*HDCISRKP HRLRPKSP*AGKQLQQSLRIQN QCTKITSIFIHQ*QANRKSNHE* TPIHNCFKENKIPRNPPYKGCEG PLQGELQTTAQ*NKRGYKQME EHSMLMGRKNQYRENGHTAQ

SEO ID	SEQ ID NO:	Met	SEO ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
]				sequence		
3885	34253	A	3925	1	1251	MKAEINMFFETNENKYTVYQN
5005	3 1233	` `	3,23	[123.	LWDTFKAVCRGKFIALNAHKR
						KQERSAMNTLTSQLKELEKQE
						KTNSKANRKQETTKIRAELKEI
1						ETQKTHQKINESRSLFFEKTNKI
						DRPLARLVKKKREKNQIDAIKN
						DTGDITTDPTEIQTTIREYYKCL
	1	ŀ				YANKLEYLEEMDKFLDTYTLQ
						RLNQEKVESLHRPITGSEIEAIIN
						SLPT/KKSPGPDRFTAQFYQRY\
					}	DGMYLKIIRAIYDKPTANIMLN
		1				GOKLEAFPLKTGTROGCPLSRL
	1	1				LFNIVLEVLARAVRQEKEINGIH
						LGKQEVKLSLFADGMIVYLENP
		1				IVSAQNLLKLISKFSKVSGYKIN
						VQKSQAFLYTNNRQTESQIMSE
						LPFTITTKRIKYLGIQLARDVKD
						LFKENYKPLLNKIKEDPNKWK
						NIPCSWIGRINIMKMAILPK
3886	34254	Α	3926	1	1203	
3887	34255	Α	3927	1	1233	
3888	34256	Α	3928	1	951	MKREKNQIDAIKNDKGDITTDP
						TEIQTTIREYYKPLYTNKLENLE
		ļ				EMDKFPDTYTLPRLNQEEVESL
						NRPITGFEIEA/INSLPTK*SPGAE
						GFTAEFYQSVGSSGQGNQARE
		1				RNKGYSTRKRGTQIVPVCRWH
				1		DCIFRKLHGLSPKSP*ADKQLQ
						QSLRIQNQCAKITSIPIHQ*QTYR
						EPNHE*TPIHNCYKENKIPRNTT
						YKGCERPFQGELQTTAQ*NKRR
			1			HKQMEEHSMLMDRKNQYREN
						GHTAQGHL*IQCHPHQATNYFL
						HRIGKNYFKLHMEPNKSLHSQ
						DNPKQKEQSWRHHAT*LQTILQ
						GYSHQNSI
3889	34257	Α	3929	1	814	

SEQ ID NO:	SEQ ID NO: of peptide	ı	SEQ ID NO: in USSN	Nucleotide location of first		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
3890	34258	Α	3930	1	1545	HQLKVQFRIIISIRSSCDGSAWG
						VAPTFKTRGARSRSRAAIRLGA
						ADLDEVKSSLVNESENQSSSSD
						SEAERRPQPVRDTFQKPRDYFA
						EVRRPQDSAFFKGPPYPGYPFL
						MIPDLSSPYLSNGPLSPGGARTN
		ļ				SSPAPKETWICARFGGVSLSFLE
						IGSRVLLLGRDVNRSSSLLPAQI
						PIACHFAVDGGNFIRGKGAYLL
						TFDLFGNWGLFFLIEIAVWELS
						AHSSGQSEDALELSRGTCSSSL
1						QLCWTAKALVGKGLDGGPVC
						KNSGICSTRTKTQEQMSFMEAL
						YQEGFLRETVVQAVRKVPQTP
						RKAVLEVLARAISQEKEIKGIQL
İ	;					GKEEVKLSLFADDMTVYLENPI
						VSAQNLLKLISNFSKVSGYKIN
						VQKSQAFPYTNNRQTESQIMSE
						LPFTITTKRIKYLGIQFTKDVKG
						LFKENYKPLLNEIKEDTNKWK
						NIPCSWIGRINIVKMAILPKVIY
						RFNAIPIKLPLTFFTELEKTTLNF
						IWNQKSR\IGKKILSKKNKAGGI
3891	34259	Α	3931	693	1464	ARAEVKLSLFADDMIVYLENPII
						*ARAEVKLSLFADDMIVYLENP
						IISAQNLLKLISKFSKVSRYKINV
						QKSQAFLYTNNRQTESQIMSEL
						PFTIATKRIKYLGIQLTRDVKDL
						FKENYKPLLNEIKEDTNKWKNI
						PCSWIGRINIVKMAILPKVIYRF
	İ					SAIPIKLPMTFFTELEKKNWLAI
						CRKLKLDFFFIPYTKINSRWIKD
						LNVRPKTMKTLEESLGNTIQDI
						GIGKDFMTKTPKAMATK/DQKS
						FCTAKETTIRVNRQPTEWEKIF
						AIYPSDKGLIS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
3892	34260	A	3932	211	2519	ENRKSNCLCLQMA*LYI*KIPSS
						QPKISLS**ANLAKSQDTKSMC
						KNHKHSYTLITDKQRAKS*VNS
						HSQLLQRE*NT*ESNLQGM*RT
						SSRRTTNHCSTK*KRTQTNGRT
						FHAHG*EESIS*KWPYCPSKC/K
						RTEIITNSPSDHSTNKLELRIKKL
						TQNHTITWKLNNLLLNDSWVN
			•			NEIKAEIKKFFETNENKKTTYQ
						NLWDTAKAVLRGKFIALNAHI
					i	GNLERSKIYTLISQLKEPERQEQ
						TNPKASRRQEITKIRAELKEIET
						QKTLQKINESRSWFFENIKIDRQ
						LARLIKKKREKNQIDTITNNKG
						DITTDPIEIQTTIREYYKYLYAN
						KLENLEEMDKFLDTYTLPRLNQ
						EEVESLNTPITGSEIKAIINSLPT
						KRSPGPDRSTAE/FYHRYKEEL
						VLFLLKLFQSTEKEG\GRDTTK
						KENFRPISLMNIDAKILNKILAN
						RIQQHIKKLIYHDQVGFIPGMQ
						GWFNICKSINVIFQYTNNRQTES
						QIMSELPFTIASKRIKYLGIQLTR
						DVKDLFKENYKPVLNEIRGHK
						QMEEHSMLMDRKNQYCENGH
i						TAQGNL*IQCHPHQATNDFLHR
						IGKEEVKLSLFADDMIVYLENPI
						ISAQNLLKLISNFSKVSAYKINV
						QKSQAFRYTNNRQTESQIMSEL
						PFTIASKRIKYLGIQLTRDVKDL
						FKENYRPLLNEIKEDTNKWKNI
						PCSWVGRINIVIMAILSKVIYRF
3893	34261	A	3933	2	1304	
3894	34262	В	3934	141	2008	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
3895	34263	Α	3935	1	1845	MVISVDAEKTFNKIQQPFTLKT
			:			LNKLGIDGSYLKIIRAIYDQPTA
						NIILNGQKLEAFPLKTGRRQGC
		l				PLSPLLFNIVLEVLARAIRQEEEI
					:	KGIQLGKEEVKLSLFADEMIVY
						LENPIVSVQNLLKPIRNFSKVLG
			•			YKINVQKSQAFLYTINRHTESQI
		İ				MSELPFTIATKRMKYLGIQVTR
						YVKDLFKENYKPLLNEVKEDT
						NKWKNIPCPWIGRIN\ILKMAIL
						P/KELEKTTLKFIWNQKRACIAK
						SILSKKNKAGGITLPDFKLYYK
						ATVTKTAWYWYQNRDIDQWN
						RTEPSDIIPHIYNHLIFDKPDKNK
		ĺ				KWGMGSLFNKWCWENWLAIC
		ĺ				RKLKLDPFLTPYTKINSRWIKD
	ļ					LNVRPKTIKTLEENLGNTIQDID
						MGKDFMSKTPKAMATKAKIDK
						WDLTKLRSFCTAKETTIRVNRQ
						PKEWEKIFAIYSSDKGLISRIYK
						ELNFTRK\NNPIKKWAKDMNR
						YF*KEDIYAANRHMKKCSSSLA
						IREMQIKTTMR/YHLTPVRMAII
			:	-		KKSGNNRTRENYFKIHMESKKS
		ļ	:	: -		QNSQGNRKEKEQSWRHHATRL
:						QTIVQGYTVAKTACYWYKNRP
<u> </u>						TDQSNRTENQEIRLHTYNHLIF
						DKPDKSNGETTPYSINGARITG
3896	34264	Α	3936	1	700	
3897	34265	Α	3937	1	3489	MKSGHPEKEQDNSDVQETREIT
						IRGLLCTALMRHSTGAIAYLGV
						LSGSASLKLAGVPLRCCEGDKD
						AGHPLETQTALCERGRGARSLV
	İ					GNTIMTSQPVPNETIIVLPSNVIN
						FSQAEKPEPTNQGQDSLKKHLH
						AEIKVIGVNLIQNVLERGWGKC
						QEMIYVLGLDICRPFFVSRVSEE
ŀ						GRMGQRGEEDANSLDFPPASLL
						CLICQEQGVNGESCSPVGMYH
						REIVPVYEVLSVITGLQIQVFSG
						KEADSVIKRS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	i .	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
3898	34266	I _A	3938	120	1331	 TSLGPIYCAVRHTSSCRKVGSSR
3696	34200	^	13730	120	1331	CQHGCQLSVRLQLDQAHHKQL
						PWLAPKNAVVPKSLEDPQGRK
		l				EGVTALTPEAPRSGPPKRLQLFP
						PSLCPNCSKQGAYFSPFGVTAT
						LLATPFSRSRVLVLQPGRMRYA
						`
						DKWRVSKMKRCCIEQWNSSEK
						THSARASHSLQRQRGGPRVGGS
			ļ			SLQAGCHVISAALSKEEALEWV
						ASFCRQIVQSYLRPLLCSGADP
						GAFMOLRGEELRSLELSLSYTP
						PSNEFKISMKLEAQDPRNTTST
						CIATVVGLTGARLRLDGSDN
						KNDFWRLVDSAEIQPIGNCEKN
						GGMLQPPLGDSFHCDVRVSILD
						LFCFLLSELPFTIDTKRIKYLGIQ
						LTKDVKDLFKENYKPLLNE/IK/
ļ						EDTNKWKNIPRSRIG*INIVKMA
2000	24267	ļ	2020	1	1421	ILPKDFG
3899	34267	Α	3939	1	1421	MDSMSGGGQYRKINGNPTSVK CPLLLLPAILTPEPVNRWRQSC
						KAFARHSPLAFRVTISTSTFFDG
		1				
			ŀ			LLVTGLYTSTSVQASQSIGGSSA
	·	1				FGFVLEVLARAIRQEKEIKGIQL
		1				GKEEIKLSLFAGDMIVYLENPIV
		1				SAQNLLKLISNFSKVSGYKINV
						QKSQAFLYTNNRQTESQIMSEL
		1				PFTIASKRIKYLGIQLTRDVKDL FK\ENYKPLLKEIKEDTNKWKN
		1		:		IPCSWVGRINIVKMAILPKNWK
		l				KLKFIWNQKRAHIAKSILSQKN
		l				KAGGITLPDFKLYYEATVTKTA
						WYWYRNRDIDQWNTTEPSEIM PHIYNYLIFDKPEKNKKWGKDS
		l				!
						LFNKWCWESWLAICRKLKLDP
						FLTPYTKINSRWIKDLNVRPKTI
						KTLEENLGITIQDIGMGKDFMS
}						KTPKAMATKAKIDKWDLIKLK
						SFCTAKETTIRVNRQTTKWEKI
						FATYSSDKGLISRICNELKQIYK
			<u> </u>	<u> </u>		KKTNNPIKK

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
3900	34268	A	3940	3	1566	IQTTIGEYYKHLYTNKLENLEE MDKFLDTYTLPRLNQEEGESLK RPMAGSEIEAIINSLPTKNSPGP DRFTAEFYQRYKEELLISNFSK VS/VIQNQWEKITSIPIHQ*QTNR EPNHE*TPIHNCFKENKI\LGIQL TRDVKDLFKENYKPLLSEIKED TNKWKNIPCSWIGRTNIVKMAI LPKDKTSKYIDVDENEGSHCGK RKYKYGMEKALEILARAIRQEK EIKGIQLGKEEVKLSLFADDMI VHLENPIISAQNLLKLISNFSKV SGHKINVQKSQTFLYTNNRQTE SQIMSGLPFKIATKRIKYLGIQL TRDVRDLFKENYKPLLNETKED TNKWKKNILSSWIGRINIVKMA ILPKVIYRFNAILINLPMTFFTEL EKTTLKFIWNQKRACIAKTILSQ RNKAGGITLRDFKPYYKATETK TASEMKYYLENKIPFKVLHMV YNVPTHPPFIGDLHPNTKVVSL PPNITSLIEPMNQGVISAFKDCY LRKTFVQAVATPEGETEMTVM OFWKDYNT

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SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
3901	34269	A	3941		2580	MVKGSIQQEELTILNRYAPNTG APRSIKQVLSDLQRDLDSHTIIM GDFNTPLSTLDRSTRQKVSKDI QELNSVLHQADLIDIYRTLHPK STECTFFSAPHRTYSKIDHIVGS KALLSKYKRTEIITNCCSDHSAT KLELRINKLTQNRSTTWKLNNL LLNDYWVHNEMKAQIKMFFET NENKDTAYQNLWDTFKAMCR GKFIALNAHKRKQERSKIDTLT SRLKELEKQEQTLHSKDSRRQE INAEKAFDKIQQPFMLKTLNTL DIDETYLKIIRAIYDKPTVNIILN GQKLEVFPLKTGTRQGCRLSPL LFNIMLEVLARAIRQEKEIKGIQ LGKEEVKLSLFADDMIVYLENP IISAQNLLKLISNFSKVSGYKIN VQESQAFLYTNNRQTESQIMSE LP\FTIASKRIKYLGIQLTRDVKD LFKENYKPLLKEIKEDTNKWK NIP\CSWVGRINIVKMAILPKVI YRFNAISNKLPMTFFTELEKTTL K\FI*KQKRACIAKSILSQKNKA GGITLPDFKLY\YYKAIVTKTA WYWYQNRDIDQWNRTEPSEIIP HIYNHLIFDKPDKNKKWGNDS
						LFNKRCWENWLAICRKLKLDP FLTPYTKINS\RWIKDLHVRPKT IKTLEENLGNTIQDIGMGKDFM TKTPKAMATKS\KIDKWDLIKL KSFCTAKETT\IRVNRQPTEWK KIFTIYPSDKGLISRIYKEPKQIY

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SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/340,217	sequence	or peprior sequence	determin, (-possible nacreorde insertion)
3902	34270	A	3943	5	2130	QRLRRQHRLEQKTRSTTCHMR
		1				QSTTRSAERADTRIESMPT*TTP
						*HTDDVP/SHCTNSTHTATTSPS
						THNHQQTLTRVANVAQIRRTDI
						SSIAATEWISTINTHNYACRTRA
						VSQRRYVSDFEKRERPTSNTPE
						PRFRLVVSTPTLVGTTQGTYQP
						PPARSRISIASHLLPPLSLLPSSL
						DLDGRSTLACSSSVSQFSGSRGS
						PSSYIVATTRVISDVDYDMTTY
	1					YNSTITPIS/PSARS*CQRSVHLL
						LSSHRTYRHPLVSRHTSQEHSL
İ						GGPLHRH*YNPVGSRAAAWAS
1						KSALV/SVSLEALVVSALI*LVA
						TRQRLVGICRTTPIRARSSVVR*
ŀ						VTRYQPNQRAPLIHATYHLLDR
						GQHPQSMQTISHWTTPPCWCL
						VCGKKSSLPCCSTSSMSTRRNQ
						YDTLSLTTSWVL*SSIFWLAFIL
						LPRTSLPWPTVS*LAANA\SSGS
						TPVNSSFRT\SVRRSKLVVPANE
		1				IETPSFVVVTKFSRSASSYDCSIE
						YASTYAINITIVNSYVFA/PTHTT
						REHTISYALTSPGQPQNKTRIPE
						LQWAF*AVRPSTQ/PSTVIYHAP
						TSQAIASCALHSLLGCLLGSAT
						APLPPTWTTPPPPPQLRTT*STG
						SLPHPPSC*TRP*PLAPRN*PFTG
						MSSQHCIPT*PQLASHSIALRG/S
						RARPTTSQTSIAS/SHSHS*LSHV
						Q*RPLSDQRSPLDHHAHSSILYA
2002	0.071	<u> </u>	2044	254	004	RASRISCLRVCAV
3903	34271	Α	3944	254	884	MTPNYTSRLFLHMGVLFYPFYR
		1				RLT*HIRTHINLKGWK/NRHFM
		ŀ				QMDTKNAEKALDKIQHRFMIK
						TLSKISIQGTHF\KIIKAIYGKSTT
						NTILNGEKLKAFPLRTGIKQGCP
						LLPLPFNIVLELLARAIRQEKEIK
						GIQIGKEEVKLTLFADDMIIYLE NSKDSSRKLPELIKEFSKVSRYK
						i
						VNLHKSIALLYTNSDQAENQIK NSTSFTI
3904	34272	В	3945	52	843	INGLOCAL TENE
3704	137214	T <u>n</u>	2743	J ²	1073	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
3905	34273	Α	3946	1	831	MEMLYCLIAEAGHISSRMATNT
ĺ						ANSAGLKPTCVCVALPLPPGRS
						APHRSCSQAGRELPGQGPRYYR
						HLPQLSILHSIGEGQGCGFWSER
					İ	SFKGYPERPAGAAGVCRLQGC
						GRRGRGAPFRTTDFSSRPRGAA
						ERADQGPRAGSPWPRTTSGAQ
						RGRAQGQGHTARRRGNSNPGP
						SRARQASRRRRPATSGPPRGSP
				}		RPDRPRRRSPFYRSSSRETSRPP
		1				EGPRRPRAPALSAPAPGQPARP
						RPREPVPCGAVFTARDRLRPPA
						ATSHAPFSAANPRR*HRPGGPG
						ARRLGDAQLSRRST/SGAPRCS
		•				QTRSR*PTCVCVALPLPPGRSAP
						HRSCSQAGRELPGQGPRYYRHL
						PQLSILHSIGEGQGCGFWSERSF
					1	KGYPERPAGAAGVCRLQGCGR
						RGRGAPFRTTDFSSRPRGAAER
						ADQGPRAGSPWPRTTSGAQRG
]				RAQGQGHTARRRGNSNPGPSR
						ARQASRRRRPATSGPPRGSPRP
						DRPRRRSPFYRSSSRETSRPPEG
						PRRPRAPALSAPAPGQPARPRP
			,			REPVPCGAVFTARDRLRPPAAT
						SHAPFSAANPRR
3906	34274	В	3947	250	281	
3907	34275	Α	3948	3	639	DHTCRLRQRLRLRVLVGPVPG
						AGPAG*KGCYGGRSANHHGAP
						ASCHLARSSCGPRLPGRYSAQQ
						PRARCAASGLCGWTAPAADPV
]						PSEVLASQEVQLLCAGE*SGSC
						GPTHADLQPSPGGTGEDGAAR
						AKRDLPGSVGERAAAPASGRL
1		1				RACPGRPAGAPGPRARPPGGTA
						ALAQPPRPQGAAARPPSGIGWP\
					}	GNNGSAQSKGRALMEQAAG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
3908	34276	I A	3949	161	2377	
3700	34270	` `		107	23 / /	CTGYQLFKPSLISWLEEEEELST
						LPRVLQEWKMCLKTKGPALW
						QDNFCLKTLNGIQLARNQNGEE
						LYDCKQCEDVFCKHPCLKTNM
						STQNRGNTSECIQYAKDLLSLY
						NKTSTIRKVSVFSKHGKSFRL\F
		l				*MFRSRESVHKINPLK/CTDYGK
						AFIYQSYLEAHRKTQSGEKLNE
						WKQCGEAFTHSTSHAVNVETH
						IIKNPYECKECGKDFRYPTHLN
						NHMQTHIGIKPYKCKHCGKTFT
						VPSGFLEHVRTHTGEKPYGCKE
						CGKAFGTSAGLIEHIRCHAREK
						TFKCDHCGKAFISYPSLFGHLR
						VHNGEKPYEHKEYGKAFGTSS
						GVIEDRRSNTGQKRFDCDQCG
		ļ				KVFVSFSSLFAHLRTHTGEKPF
						KCYKCGKPFTSSACLRIHMRTH
						TEERLYQCKKCGKAFTKCSYLT
					!	KHLRTHAGEKPYECMKCGKAF
						TERSYLTKHLRRHSGEKPYECK KCGKAFTERSDLTKHLRRHTG
]					
		i				DKPYEYKDCGKAFVVSSSLVD
						HLRTHTGYKPYKCNACEKAYS
						RSCVLTQHLKTHAAEKTSECN
						ACGNSFRNSMCFHDRLKTLTKI
						KPYKCKDCGKAFTCHSDLTNH
						VRIHTGEKPYKCKECGKAFRTS
		l				SGRIQHLRTHMGEKPFECDQCG
· ·						KAFA\FSQLVLHI*KHTREKPCG
		_	 			CEECGKTFAVSSSLTEHVKIHR
3909	34277	Α	3950	6	455	GLLHERQAEARCSICLDYLRHP
ľ		1				MTTDCRHYI*SARIHQCW*ELQ
						DISPCPVCLQHCPDKNLKRNFQ
						LCHMTDIAKQLLTTARRKRKL
						QGEEPVCRKSDVALFCEKDPEL
						LCHQYRVSLDH*DH/SPMPIEQ
		<u> </u>				AAAKHRKQFESYIEPLEKQV

SEQ ID		1	SEQ ID NO:		ł .	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
3910	34278	A	3951	2	1009	 WNGHRMN*MQSSNGLEWN/QS
	12,0	` `		[,	SNGKEWNHRIESNGIIIAWN\Q
		ŀ				WYQHQTEKNGIFEWNRRESSN
						GPEWNHVMEWNG/DNPWTRM
						QSSSNGIEWNHRMDSNGIIFQW
						NGNGNHRIGIEWNYDQ\SNEWI\
						QWNQHQTEKNGIIKWNRRESS
1						NGPEWNHLMEWN/ENNPWTR
						MQSSSNGIEWNRMESSNGLEW
						NNH*TESNGSVESSSDGNER/QS
						SSNGIAWNHHKMESNGINIKW
ļ						NQMESWN/WN*MNRMELSSNG
						IEWNQHQTEKNGIIEWNRRESS
						NGPEWNHLMEWN/ENNPWTR
						MQSSS/NWNRMESSNGLEWNN/
						QLNGIEWNHHRMEMNGIIIEW
						NRIELWN
3911	34279	Α	3952	1	1494	MASLLGAPRLAGWASGAGALS
						RGWAIRPADTGGNLGPVPRVPL
			<u> </u>			PPDPVLTARWAPGVNSGSQFSC
						HCQAPILEMGHKGSSPGLGDAE
Ī			:			VRAITVQCIRPIDGPQQPPGGGS
						AGRRLTIPASTQEWAQLPVGRV
						LANVLTEGGDTGNQPIPQRSLC
						RPQPCSHAETWGEVEAQVPAQ
						SNREQPAAAPGCGPGRGETGA
						RPETTFSPRRAPPNPYDEEGVR
						WSLEFMLCGTDGPVQPVQHQE
1						GPAARLQLIRGGSLILESEGTLR
		ŀ				G/SPVLQTDQPASHLLHTQGFW
						A/AALSAVCL/HQNIIHSGSALL
						APATRAAWEQIQRSEGGTAQL
						LRRLEGYFSNVARNVQWTYLQ
						PFVIVTTNMILAVDIFDKFNFTG
						ARVPWFDAIHEAFPRELESSISF
İ						PANFFKPPEEKEGPLVRPASRK
						TTPQTTRPGPGTEREAPISR*KR
						HPDDTG*FTFTLGIVYCTPGQLP
						PEPYDPNRRSLWLPHWPIINTS
						MVSALVYSEGAPLPSPL
3912	34280	Α	3953	1	681	MGQLLDKNTPSHGARTREECG
						RERLCVSPSQTGDTPRTSAYLC
						VGGPAWSPLSESRPAGSSGCPW
					l i	IKPPDPRYSPIGLCSLLTTEMMS
				:	į l	RQPRTDLRGQTNPAA\PSAPVPL
						SCSQNLPVWPSLMAGTTWHSP
						L\SPSCFWHSPGHN*H*CCVSKD
]						*KSLFWEPTA/YSPLLPSTSP/SS
						KSMQPPKPRSNADSSVQASLIP
						RAMSSPTVSPWIMGNGSQGFD
						HIAVSMWDD

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
3913	34281	ĪA	3954	136	420	RESRNLSRGEESEDPAEPS\RNG
	13 1201	ļ.,		130		*EHPLCYLQ*EFLQVLTMLQAE
						GTMHHFRSICQVNRNFLERGH/
		1				SPPSPAPPPETHTGSPRPPSGRSR
		İ				IRAYLH
3914	34282	A	3955	1	1782	THE TENT
3915	34283	В	3956	1	3070	
3916	34284	A	3957	104	279	STTHPSVHE/QEEEEEEEEE
		<u> </u>				EEEEEEEEEEEEEEEKKEC
						SKAQCKHFPLSEVL
3917	34285	A	3958	1	252	MTVCIHIASEDLPVGRDVEVED
	3 1203	ľ.	3,50	l'	232	SDIHDRDPGLGDKSETPSE/EKK
						EEEEEEEEEEEEEEEEEE
						EEEEEEEEEEEEEEEE
						NFLQHYLHL
3918	34286	A	3959	2	368	MEQMENE
3919	34287	A	3960	239	432	CLWLFQEEEEEEEEEED*EEE
	34207	^	3700	237	132	EEE/EEEEEEEEEEEEEEEE
]				EEEEEEEKIFLGHRVGI
3920	34288	A	3961	1	577	MQIPSLHKLKKEEEEEEEEEE
3920	34200	^	3901	[*		ERRRIGRGREKKEEEEEEEEE
				İ		EEEEEEEE\EKKKKEEKKKKEE
						EEEERRRKKKKERKEEEEEE/G
]	1				KEEEEGEEGEEEEEEEERRK
	1	:				EEEEEEEGEEEEEESCLMGP
						MCVHIHP\DKDLYSLGPPAQRF
						TGSHAELPT*KARRSSSWTAAS
2021	24200	 	2062	227	550	RGCAARDPPRRCSPA
3921	34289	Α	3962	327	559	PKGRTPSPSCIHRYPCQTPRPHE
						P*GCHCPEEK/PRPRVWGPSRC
						MPLGSVQEKRPCPAPGGVQGSF
2022	124200	ļ	2062	1	577	RVSPLMMLTRL
3922	34290	A	3963	1	577	MQIPSLHKLKKEEEEEEEEE
						ERRRIGRGREKKEEEEEEEE
						EEEEEEEE\EKKKKEEKKKKEE
						EEEERRRKKKKERKEEEEEE/G
	1					KEEEEGEEGEEEEEERRK
]						EEEEEEEGEEEEEESCLMGP
						MCVHIHP\DKDLYSLGPPAQRF
						TGSHAELPT*KARRSSSWTAAS
						RGCAARDPPRRCSPA
3923	34291	Α	3964	157	272	WCNGSPLYSGW*LVGMESLGR
						MHKDLWTRQPNQDQDLQ
3924	34292	В	3965	1	3723	
3925	34293	В	3966	1	573	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
3926	34294	A	3967	3	424	AGGQALQGPGQGRGSQGVVG/ PGTGSSGAQHLGKHYPVLSGGS ERSWGRSHTPAGAGC*VRGRR AGGRPGTGHSRPPGGSCLSPAP PNSARWLGLGAPWQAGAGLR DPGDWRRGQGGPGWAWCPGQ PPAQPHTCPPNSTARY
3927	34295	A	3968	3	1238	RGAERRAWSRGPACTRRGPAD WAAAGAGRPCPQRRGVCCTAA VPGAARLSCPTGPGPGDPGRRS LTGQGS*GLGAAFGGWTGALP SWHS*SQGWQTDPVR*VRGTE RDICTGL*QPCPPGGLQTGSGG LEHSLPWPGIGIQAP*GPNHPCR LPRS*ALSAGG\SGGQALQGPG QGRGSQGVVG/PGTGSSGAQH HCCPPYTPPG*HPIPSLLALGPQ SLQPEWAHSGTASGEQHSAGE HGMGTTGH*LPGLCSRCVLGK HYPVLSGGSERSWGRSHTPAG AGC*VRGRRAGGRPGTGHSRPP GGSCLSPAPPNSARWLGLGAP WQAGAGLRDPGDWRRGQGGP GWAWCPGQPPAQPHTCPPAGS LPGAAPGVLCAA*GPAAGV*A
3928	34296	A	3969	3	415	ETGRHRSQQSVSSPPVQPRGKR AMYHSAAELVSRGFPRPPVQAP AEPAGAAEGVHSQPASRQEA/G S/TEVRGQAHRFVSPPNAAGAG DG/PDPQSLLAPTNRPCPPGGISP ARSEPVPPAPGRAAP*CFPDLPG LAPPLC
3929	34297	В	3970	1	657	
3930	34298	A	3971	125	524	EAEALENQSQPCDTG/PQSAFSP PGSTQHPRSQLSQCKQRYQDLQ EKLLLSEATVFAQANELEK*RV ILS\GEPLLKQDSKQVQVDLQD LGYETCGQSKNEAEQEETTSPE HEEHSSRKEMVLVEGLCSEQG
3931	34299	A	3972	1		MGQVWGLVHFTLEVFHTGDEE EQEYSEVTEDVTEHVYLPAKA KVAKEEEAGIQQARQEGDLEA WQFPVRIHPPDQQENITATFEPF PFKLLKELKQAINQYGPGSPFV MGLLKNVTVSSQMIPTDGDPLT RACLTPAQFLQFKTWWADEAS IQAARNAWAQPQINITADQLLG VGGWAGLDAQFVMQDDAIEQ LRGVCIRAREK\IT*CGEQYPSF

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
3932	34300	Α	3973	350	1078	GSNRRSNRAEQLRGVCIRAWE
						K\ITSGEEQYPSFSIVKQGPREPY
		1				TDFIAWLQESLKKVIADSAAQD
		l				IVLWLLAFDNANPDCQAALRPI
						RGKAHLVDYIKACDGIRGPAG
						RLSWLSLQGRPAGAARCTSSSS
				į		DWKAQLAPGPHAGPLRCPAFR
						ARQPRLPAVLSAGAAFRDCPLS
						CVQALLGGSAGPGDRLPGHYS
						ALVALSLPFVKEATMNRWSRN
						HRSAFFLFSANAHGAEGVLSHT
						VASR
3933	34301	A	3974	2	630	WDNCGLWFIPSWNLFTLMMRR
						KESLMK*QKK*QSRFVCQLKLK
						/PAKEGEVYPYPSAPPPYFEEKE
						WPDPPDLSFLEDAGQKVIAPVT
						VQAAPQAIALSSIQAGIQQARR
						EGDLEAWQFPIRIHPPDQQGNII
						ATFEPFPFKLLKESKQAINQYG
						QGSPFVMELLKNVAVSSQMIPT
						DWDALAQACLTLTQFLQFKTL
						WADEVSIQAACNA
3934	34302	A	3975	264	634	WSSRCQHSSRPQASESWFPG*G
						PSFWPRIQGDEKTGAGGHP*LG
						C*PGMTGQGFSTKCQHTCLMW
	}	ĺ				GSHWAQEAPENAPGTSCPGSSG
						SWVLRSSLQRQKSAWSPG/ASM
						PAPKMPFLTPSSGFS
3935	34303	Α	3976	3	410	KKKVWREEKERLLKMTLEERR
						KEYLRDYIPLNSILSWKEEMKG
						QGPK*\EENTQETSQVKKSLTEK
					·	VSLYRGDI/L*VDAIVNAANASL
						LGGGGVDGCIHRAAGPCLLAE
						CRNLNGCDTGHAKITCGYDLP
						AKCEYN
3936	34304	Α	3977	74	432	MLHNLRPRTLTTRTRCPSTPS*T
	1					TT*ATPPTTTHGSAGPRAAHLR
						RTGTRRWRAPRRARSCTRSSPR
						RARAASTPPLAPARELRSPASPP
						SCEQSAAPPSGRNGGNFPESIFV
						KTINSN

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
3937	34305	Α	3978	2	894	WGGYGMRTGPRTLTTRTRCPS
						TPS*TTT*ATPPTTTHGSAGPRA
						RTCAAQRTSGSHPSQRRTSSAA
						PGV*RANVFPWAH*MKRV*TT
						LENLTA/PEMAMPPAPHVIFAT
						DDWAAMVHPSARVPGLDGTG
						ALLVPTGVCAPGPCKCPLTSSS
						VTHTRLKTTLSVPPSARQTGRC
			-			RSPSDLRCSYPPDEQPVCVPKC
			}			GSPWVSVLVAWIQSESAVLDPR
					·	HPQHPYLYPVDMQNLLTNLGE
						PPQARALAAKLLGRPSSSQSGS
						RVPAVWAQAGNATYITVHTLC
						SHNTHMSPVRVKRFTHLG
3938	34306	Α	3979	157	570	
3939	34307	Α	3980	1	936	
3940	34308	В	3981	257	3934	
3941	34309	Α	3982	210	4286	MPLKTRTALSDDPDSSTSTLGN
						MLELPGTSSSSTSQELPFCQPKK
						KSTPLKYEVGDLIWAKFKRRP
		ļ		•		WWPCRICSDPLINTHSKMKVSN
						RRPYRQYYVEAFGDPSERAWV
ļ						AGKAIVMFEGRHQFEELPVLRR
		ŀ				RGKQKEKGYRHKVPQKILSKW
						EASVGLAEQYDVPKGSKNRKCI
						PGSI\KLDSEEDMPFEDCTNDPE
						SEHDLLLNGCLKSLAFDSEHSA
						DEKEKPCA\KSRARKSSDNPKR
						T*L*KRATYNFEAH
3942	34310	C	3983	163	309	DATE OF THE PROPERTY OF THE PR
3943	34311	Α	3984	72	424	RNCGTARSQHEPLGSWLQDTP
				·		QPP*TLELAGNLPGD/F*PGPGK
						EQGMFVCHPIRQPLPRPLPGSSH
				:		QSMPTAQPPLSSSSALLPALPAG
						FPVTQGQWTKLQVQAPAPFHL
						PPQVEAV*AFYQKQMLVPCSL*
	1					SMPTAQPPLSSSSALLPALPAGF
						PVTQGQWTKLQVQAPAPFHLP
		\perp				PQVEAV
3944	34312	Α	3985	1	347	KWQRFVLTGIDTYSRYEFAYPA
1		1				CHASTKTTIHGLMEFLIHHHGIP
						HSIASDQGTHLMAKEVRQWAH
	1					AHGIHWSYHVPHHPEVAGLIER
[WNE\GLLKSQLQHQLVNRLRRE
	1			<u></u>		LQCWLG
3945	34313	IA_	3986	1	1716	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
3946	34314	A	3987	1	737	MSSVLLRLIYQLTQKTASFEGG
		l				PEQKALQQIQAAVQAALPLGPY
						DPANPMVLEVSVADRDTVWSL
						WQVPIGESQLGFWSKALPSSAD
						NYSPFERQLLACYWALVETEY
		1	ļ			LTMGHEVTMLPELPIMTWVLS
ŀ						DPSSYKPAPMASWGVPYGQLT
						EEEKTRAWFTDGSARYTETTR
						KWTAVAIQPLSRTSLKDSNEGK
						SSHQQAKNGITVLAGVIDPDYQ
		1			-	DEISL/LTPQWRCHPGSSVWRAS
						SLGISHPP
3947	34315	Α	3988	2	384	CGRSGYWHSSVATKITRLRML
						RPREGRKLPPGDIMIPLN*KLRL
						PPGS/FLLLSHQAKKGVTMLAG
				İ		VTDPDYQDEISLLHNGGTGKS
						PHISDTFYGSKVASCQNTGPEK
						QDETQAQETAVYKSQIFGS
3948	34316	Α	3989	3	1273	
3949	34317	Α	3990	3	341	GLGRRQPAGSWPERRPGPSA\R
						RSTAPRRCGQAES*TERGSQPH
						QVQGQGRWGVCMKIPSHSGKS
						PDVSEVSKSRNSIISTAVTHAVV
						APEGLKRNGGGSHLRSSRGHR
						AVIF
3950	34318	Α	3991	44	243	
3951	34319	Α	3992	40	558	LGSIQVMQAVRNAGSRFLRSW
ĺ						TWPQTAG*QMTAPSSPPPPPGL
				·		CSYSCPLSHP/SLPVTVRPWPSPS
1				į		FSSQQGRGQNA/APGPSAQALD
						SSKTLRPSRKLNRTRLPATPSSG
						EPHLDQPSGDPQPLTLARHPPES
						EPVNFQLCHLLSVGPYANKSEP
						QPSHLKMRIMLREVVRIT
3952	34320	Α	3993	335	581	RRHLFLQWGQRAWRLQVAAA
1				Ì		GTTRPTSAMGIRCSEGAAARAT
				1		AARA*TAGPEPLE/PAANPPPPL
				1		TASALRAPPSFVLPQCTR

SEQ ID	SEO ID NO-	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
2052	34321	I A	3994	216	1159	SWHGPPGANTVAAAAGPEE\K
3953	34321	A	3994	210	1139	
						AALKLRPTHGWPVRATDVHDV
				İ		ILKASGESEWRGGSRASHQMSP
				1		VAMATALGEGVPVRGPAAGSL
		ŀ				RLLPGSSAPLGRDAISSCNGVN
						GLETTGGRCRHNAPNKRHGDP
						LLEGAAAR/AQAARA*TAGPEP
						LE/PAANPPPPLTASALRAPPSF
				•		VLPQCTAAPRDPSAAGAAN*G
				1		KAQSRNC*NEPFAYGGGTHGT
						GAGAAVTVAADGN*LGSIQVM
						QAVRNAGSRFLRSWTWPQTAG
						PWPSPSF/VFPAGSWPERRPGPS
		1				AQALDSSKTLRPSRKLNRTRLP
				1		ATPSSAFTLPFQERRAL
3954	34322	Α	3995	1	738	MTKRGHGTAWAVASKSASPKP
1						WQLPHSVEPVGTEKSRIEVWEP
			}			LPRFQRMYGNTWMSGSSLLQG
						NQNLHAERYCNSTLERNDTPIE
						SLKPKRESEDGLGEHNGSTMEE
						VGAETRVQRHWVRVSMTELAL
						ASDAHMWGSNPGQRVTGVMV
						GECGTMLGDTQVLLSNPCGDR
						ARRAYSTAPDYAVCGNGGKVK
	1					LNEQRFGSTNKQGKAAYWME
		1				ALRPEPLCWQSNYPEAAAVGK
	1					PKAAYTKKLHGEDS*AIPVVTE
						LGERIAQLLIMLYV\KWGKSEIK
						RT/G/GFGSTNKQGKAAYWME
						ALRPEPLCWQSNYPEAAAVGK
		ļ				PKAAYTKKLHGEDS
3955	34323	С	3996	87	329	
3956	34324	A	3997	3	122	
3957	34325	Α	3998	1	156	
3958	34326	Α	3999	1	353	
3959	34327	Α	4000	1	201	DELLACACIONE DE LA CORRECTION DE LA CORR
3960	34328	Α	4001	56	207	EEKKEKEKEKEKEK/EEEEEEE
				1		EEEEEEEEEEEEEEE
		<u> </u>		<u> </u>		EEEEEEEEEEEEEE
3961	34329	Α	4002	1	174	MNRC*RHIYSSNEVH*KEEEEE
						EEEEEEEEEEEEEE
		_				EEEEEEEEEIPLSSL
3962	34330	Α	4003	1	278	MTSYKFTEPKNGIWQLHEAAQ
						LDTTYNKLNKKEEEEEE\EEEEE
						EEEEEE\EEGEGEEEEEEEE
		<u> </u>				EEEEEEEEEEEEEGVIL
3963	34331	Α	4004	144	429	DLPREEYALLPAGPRRRCRHTH
		1				RYEPNPEFGAKHSCPAA*HRAA
						PATSDTQE*HRSNAFGEEEEEEE
						EEEEEEEEEEEEEEEEE
			L			EEEEEEEEEETLFSNM

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
3964	34332	Α	4005	3	122	TEEEEEEEEEEEEEEE
		<u> </u>				EEEEEEEEEEEEEEEEEEEE
3965	34333	В	4006	.1	300	
3966	34334	Α	4007	1	1226	MPSSKGVHVHSPPRYLAAKDF
						KMINKELTAATFMEVIAEDNRF
						IYDGIDSNFEPELVFLEFFEALLS
						FAFICVTDQMTKSYTNVPADD
						VSGNKHETIYTILNQDAQNKSP
						SAVMSHESDAAHSDSARSSSSK
						LELSPDVNKIRKSEAMVKEKKK
	1					ADKKGEKSARSPSSLSDNLDFS
					•	KQDGNTTRQEMSPAGVPLLGM
ł						QLNEVKPKKDRQNVQQNEDAT
				:		QYEESILTKLIVESYEGEKVRGL
						YEGEGFAAFQGGCTYRVSCPFE
						NLQEGEEGRLCEECPDEPRRVH
				İ		VAGRSMYEGEVVNGMRNGFG
						MFKCSTQPVSYIGHWCNGKRH GKVGEVATWRAEKKKKEEEEE
						EEEEEEEE\EEEEEEEEEEEE
		1				EEBEEEEE/EEEEEEEEEEEE
						EEEEEEEEKIRP
3967	34335	A	4008	453	705	LLSIVQAEAVSENSHPILPRVSR
3907	34333	A	4008	433	103	SGWGQKEEEEEEEE\EEEEEEEE
						EEEEEEEEEEEEEEEEE
						EEEEEEEEEEEGRRRRSSP
						SCYSITPELSCKLGHR
3968	34336	A	4009	93	705	ESSTQTCSGFWTGCTALHRWR
		'				GMPERCPPESRDS*TRFPQSSLP
						GHKT/SEKEEEENRKEEEEEEKE
						KEK/EEEEEEEEEEEEEEE
						EEEEKEEEEEEEEEQEEEEDDEE
			İ			EEEEKSCSVNVSLIELPWDPKA
1						YSRLAPLSSQPGPAVKVPTEHLI
			ŀ			AKLEDCVQGFTYLTVEKRWAR
						AVTGAQELGVDYPRNEKCKPH
					:	NNGYDND
3969	34337	Α	4010	1	3189	
3970	34338	Α	4011	1	5127	
3971	34339	A	4012	209	3816	QGRPTFRFRKYREHHKDTPREE
						QLQDT*SSDSPKLK*RKKC*GQ
						PERKVKLPTKGSPSD*KRISRQ/
						KTLQARRQSWFFEKINKIDRPQ
				1		ARLIKKKREKNQIDTIKNDKGD
						ITTDPTEIQITIREYYKHLYANK
						LENLEEMDKFLDTYTLPRLNQE
						EVESVNRPITGSEIEAITNSLPTK
						KSPGPDGFTAEFYQRYKEELVP
						FLLKLFQPIEKEGILPNSFYEASII
		}				LIPKPGRDTTKKGNFRPISLMNI
L ,_						DAKIL

SEQ ID	1 -		SEQ ID NO:		1	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN 09/540,217	location of first codon for peptide	of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
ŀ	sequence		09/340,217	sequence	or peptide sequence	deterior, (-possible nucleotide inscritori)
				Joquent.		
3972	34340	В	4013	1	3570	
3973	34341	Α	4014	1	2347	MELKTKARELHDECTSLSSRFD
						QLEERVSVMEDEMNEMNLPTK
						KSPGPDGFTAEFYQRYKEELVP
						FLLKLFQSIEKEGILPNSFYEPSII
						LIAKPGRDTTKKENFRPISLMNI
						NAKILNKMLANQIQQHIKKLIH
						HDQVGFIPGMQGWFNIRKSINV
						IQHINRTKDKNHMIISIDAEKAF
						DKIQQHFMLKTLNKLVLEVLA
						RAIRQEKEIKGIQLGKEEVKVSL
						FADDMIVYLENPTVSAQNLLKL
	ŀ					IGNFSKVSGYKINVQKSQAFLY
						TNNRQTERQIMSELPFTIASKRI
					!	KYLGIQLTRDVKDLFKENNKPL
						LKEVKEDTNEWKNIPCSWVGRI
l						NIVKMAILPKVIYRFNAIPIKLP
						MTFFTELEKTTLKFIWNQKRAC
ļ				1		IAKSIFSQKNKAGGITLPDFKLY
						YKATVTKTAWYWYQNRDIAQ
						WNRTEPSEIMLHIYNYLIFDKPE
						KNKQWGKDSLFNKWCWENWL
						AICRKVKLDPFLTPYTKMNSR
						WIKDLNVRPKTIKTLEENLGITI
]						QDIGVGKDFMSKTPKAMATKA
						1 -
						KIDKWDLIKLKSFCTAKETTIRV
						NRQPTTWEKIFATYSSDKGLISR
		·			1	IYNELKQIYKKKTNNPIKKWAK DVNRHFSKEDIYAAKKHMKKC
						i
						SSSLAIREMQIKTTMRYHLTPV
İ						RMAIIKKSGNNRKIQ/GGIWCD
						RIL*R*TTCRVAKEIQSL*RRI/W
2071	2 12 12	ļ	1015		5050	KRLQRTLSIPVLDAV*PPMF*AS
3974	34342	A	4015	1	5073	
3975	34343	Α	4016	1	3297	MEN MEN A DEL DEFONOL DODOR
3976	34344	Α	4017	l	3514	MELKTKARELREECRSLRSRCD
						QLEERVSAMEDEMNEMKREG
						KFREKRIKRNEQSLQEIWDYVK
						RPNLRLIGVPESDVENGTKLEN
		 .				TLQDIIQENFPNLARQANIQIQEI
						QRTPQRYSLRRATPRHIIVRFTK
						VEMKEKMLRAAREKDRSTRQK
[VNKDTQELNSALHQADLIDIYR
						TLHPKSTEYTFFSAPHHTYSKT
						DHIVGSKALLSKCKRTEIITNYL
						SDHSAIKLELRIKNLTKSRSTTW
					L	KLNNLLLNDYW

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
3977	34345	A	4018	1	2666	MVKGSIQQEELTILNIYAPNTG
3711	34343	^	1010	1	2000	APRFIKQVLSDLQRDLDSHTLI
						MEDFNTPLSTLDRSTRQKVNK
						NTQELNSALHQADLIDIYRTLH
						PKSTEYTFFSAPHHTYSKIDHIV
						GSKALLSKCKRTEIITNYLSDHS
						AIKLELRIKNLTQSRSTTWKLN
		l		İ		NLLLNDYWVHNEMKAEIKMFF
						ETNENKDTTYQNLWDAFKAVC
						RGKFIALNAYKRKQERSKIDTL
						TSQLKELEKQEQTHSKASRRQE
					:	ITKIRAELKEIETQKTLQKINESR
		1				SWFFERINKIDRPLARLIKKKRE
		1				KNQIDTIKNDKGDITTDPTEIQT
						TIRESYKHLYANKLENLEEMDT
ŀ						FLDTYTLPRLNQEEVESLNRPIT
						GSEIVAIINSLPTKKSPGPDGFTA
						EFY/PESYL*QTHRQYHTEWAK
						TASIPFENWHKTGMPSLTTPIQH
						SVGSSGQGNQPGEGNKGYSIRK
		1				RGSQIVPVCRRHDCLSRKPHRL
						SPKSP*ADKQLQQSLRIQNQCT
						KITSILIHQQQTNREPNHE*TPIH
İ						NCFKENKIPRNPTYKGCEGPLQ
		İ				GELQTTAQGNKRGHKQMEEHS
		İ				MLMGRKNQYRENGHTAQGNL
						QIQCHPHQATNDFLHRIGKNYF
		İ				KVHMEPKKSPHRQVNPKPKEQ
						SWRHHTT*LQTILQGYSNQNSM
						VLVPKQRYRSMEQNRALRNNA
						AYLQLSDL*QT*EKQAMGKGFP
		1				I**MVLGKLASHM*KAETGSLP
3978	34346	Α	4019	824	3693	AWKGTTDRSTRQKVNKDTQEL
						NSALHQADLIDIYRTLHPKSTE
						YTFF/LAPHHTYSKIDHIVGSKA
						LLSKCKRTEIITNYLSDHSAIKL
						ELRIKNFTQSRSTTWKLNNLLL
						NDYWVHNEMNAEIKMFFETNE
						NKDTTYQNLWDAFKAVCRGK
]					FIALNAHKRKQERSKIDTLTSQL
					-	KELEKQEQTHSKASRRQEITKIR
						AELKEIETQKTLQKINESRSWFF
	,					ERITKSDRPLARLIKKKREKNQI
						DTIKNDKGDIT
3979	34347	В	4020	1	3765	
3980	34348	A	4021	1	4791	
3981	34349	A	4022	1	3297	
3701	12727	Ι΄,	1.022	<u> </u>	10271	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
3982	34350	A	4023		3170	MVKGSIQQEELTILNIYAPNTG APRFIKQVLSDLQRDLDSHTLI MGDFNTPLSTLDRSTRQKVNK DTQELNSALHQADLIDIYRTLH PKSTEYTFFSAPHHTYSKIDHIL GSKALLSKCKRIEIITNYLSDHS AIKLELRIKNLTQSRSTTWKLN NLLNDYWVHNEMKTEIKMFF ETNENKDTTYQNLWDAFKAVC RGKFIALNAYKRKEERSKIDTL TSQLKELEKQEQRHSKPSRRQE ITKMRAELKEIETQ
3983	34351	A .	4024	281	3030	KPRLENYMKNAEASRADAINW KKGY/LVMEDKMNEMKREGKF REKRIKRNKQSLQEIWDYVKRP NLRLISVPESDRENGTKLENTL QDIIQENFPNLARQANIQIQEIQ RTPQRYSSRRATPRHIIVRFSKV EMKEKMLRAAREKEIQTNIREY YKHRYANKLENLEEMDKFLNI YTLRRLNQEEVESLNRPIRGSEI VAIINSLPTKKSPGPDGFTAEYY QRYKEELVPFLLKLFQSIEKEGI LPNSFYEASII
3984	34352	A	4025		3290	MGELITPLSTLDRSTRQKVNKD TQELNSALHQGDLIDIYRTLHP KSTEYTFFSAPHHTYSKIDHILG SKALLSKCKRTEIITNYLSDHSA IKLELRIKNLTQNRSTTWKLNN LLLNDYWIHNEMKAEIKMFFET NENKDTTYQNLWDAFKAVCR GKFIALNAHKRKQERSKIDTLT SQLKELEKQEQTHSKASRRQEI TKIRAELKEIETQKTLQKINESR SWFFERINKIDRPLARLIKKKRE KNQIDTIKNDK
3985	34353	A	4026	1	3573	
3986 3987	34354 34355	B B	4027 4028	1	4251 3065	
3988	34356	A	4028	965	4089	TWKGTTSTSRCKIMPKYRSTRQ
						KVNKDTQELNSALHQADLIDIY RTLHTKSTEYTFF/LAPHHTYSK IDHIVGSKALLSKCKRTEIITNY LSDHSAIKLELRIKNLNQSRSTT WKLNNLLLNDYWVHNEMKAE IKMFFETNENKDTTYQNLWDA FKAVCRGKFIALNAHKRKQERS KIDTLTSQLKELEKQEQTHSKA SRRQEITKIRAELKEIETQKTLQ KINESRSWFFERINKIDRPLARLI KKKREENQID

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
3989	34357	A	4030	523	3981	
3990	34358	Α	4031	1	3429	
3991	34359	Α	4032	1	3156	
3992	34360	Α	4033	2	4943	
3993	34361	Α	4034	1	6747	
3994	34362	A	4035		3928	MAAWNLLLKSYAYWGGLRKE DFHCLDRKTLRTVSFLAALLSY ESIGGKGKLTTRKDIYTENPSV
						HHHHQRPKVDKTTKMGKKQN RKTGNSKMQSASPPPKERSSSP ATEQSWMENDFEELREEGFRRS NYSELREDIQTKGKEVENFEKN
						LEECITRITNTEKCLKELMELKT KARELREECRSLRSRCDQLEER VSAMEDEMNEMKREGKFRDK
						RIKRNEQSLQEIWDYVKRPNLR LIGVPESDVENGTKLENT
3995	34363	Α	4036	1	3638	
3996	34364	В	4037	877	8907	SNSHITILTLNVNGLNAPIKRHR LANWIKSQDPSVCCIQETHLTC RDTHRIKIKGWREIYQANGKQK KAGVAILVSDKTDFKPTKIKRD KEGHYMMVKGSIQQEELTTLNI YAPNTGAPRFIKQVLRDLQRDL DSHTLIMGDFNTPLSTLDRSTR QKVNKDIQDLNSALHQVDLIDI YRTLHPKSTEYTFFSALHHIYSK IDHIVGSKALLSKYKTTEIITNC LSDHSAIKLELRIKKLTQNRSTT WKLNNLLLN
			1	8//		OCCUCCODE*NICOCCA CDOCA LD
3998	34366	Α	4039		450	QGSPSGSRE*NSQSSAGPQCALP PAMA*VPLSWRSMGKWWKRT SCTSDST*PPSERRHWRSRKSPS AMPASFRCSSASAREMLP*KKG RCAAGSGIAPGPPETWGRTGGC PGKQATCGVSGPNANGEPVL/K YPSSSSEAHGGPGRNGRSD
3999	34367	Α	4040	2	522	
4000	34368	В	4041	102	186	
4001	34369	Α	4042	2	5417	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
4002	34370	A	4043	45	11585	KQPSGLLKFGNLILKCHPPSLTH
1002	34370	`	10.13	1.5	1303	MLSQPCA\EAPTPDPNWELA\LY
						IHPSSGIMSATVSFWSIGTA\YLE
						AQGIWEP\FRRRLS\FEASNPPFD
						VGRPFDLRRIVGISSEGNLNTLS
						CDPGHSRGFCGAGGSSSRPSAG
						SHKQ*GPSGHPHSSHSNRNSAD
						VDDVRAYNSGRTSSMTSAQAA
	Ì					SSQPANKTRPLVLDSNTGAQGH
						SAGRKSKGAKQSQHGSQHHAH
				ļ		SPLEQHPQPPLPPPVPQPQEPQP
						ERLSPAPLAHPSHPERASSARHS
						SEDSDITSLIEAMDKDFDHHDSP
					}	ALEVFTEQPPSPLPKSKGSTEGG
		1	1			PASTFTQAVDGGIQFFTDCWTE
			}			GPSSSLLAVAREVQLALCIHELL
						IHGFSQLQVSGGPGAMPDPAAH
						LPFFYGSISRAEAEEHLKLAGM
						ADGLFLVRQCLRSLGGYRQLN
						GTYAIAGGKAHCGPAELCEFYS
						RDPDGLPCNLRKPCIPPSGLEPQ
						PGSSTACETPWARPRSRPSSARP
						RRWRSSLLRRTTSGCPGTTAA
4003	34371	A	4044	1	1773	
4004	34372	A	4045	h i	663	MALWTLRPTPLLVTCMLICAPG
1007	3.3.2		1.5.12			VMGAVVAPLTILGGPLLIRAAW
						YTAGIVGGLSTVAMCAPSEKFL
						NMGAPLGVGLGLVFVSSLVDQ
1						MGRWFVAGGAAVGLGALCYY
ŀ						GLGLSNEIGAIEKAVEYWFNSF
						VCHSNQQNACSHELHDERLLG
						DMGLPILHAMLLRRLPSVDSQN
						ALSSIMLLHTALP*QSAERLFS*
						TS**EALG*YGFAYPACNASAK
						TTIRGLTECLIQHHVTPHSIASD
						QGTHFTAKEVQQWAHAHGIH
						WPYHVPHHPEAAGL
4005	34373	В	4046	147	330	
1.000	10,00	1~	1	1		L

SEQ ID	SEQ ID NO:		1	l e	•	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4006	34374	A	4047	485	1568	GEGYKADLAAATVECPICQQQ RLTLSPQYSHIPQGDQPTTW*Q VDCIGPLPSWEGQRFVLTGIDT YFRYGFAYPACNASVKTTIHGL TECLVHHHGVPHGI\ASVQGTH FMA*EVQQWAHAHGIHWSYH VPHHLEAAGLIEQWNGLLMSQ LQHQLGDNTLQGWGKVLQKG VYALNQCSIYGTVSPIARIHGSR NQGVEVAPLKITPSDPLAKCLL PFPKALHSACLEVLVPEGGTLP PGDTTTIPLNWKLRLPPRHFGL LLPLSQKAKKGATVLAGVIDPD YQDEISLLLHIGGKEEYAWNTG DPLGRLLVFPCHVIKVNGKLQQ PNPGKTANDPDPSGMKV*VTPP GKKNPRPAEVLAEGK
4007	34375	В	4048	182	662	GKKNPRPAEVLAEGK
4008	34376	A	4049	1	2250	
4009	34377	Α	4050	1	1326	
4010	34378	Α	4051	1	1614	
4011	34379	Α	4052	1	2586	
4012	34380	В	4053	1	1954	
4013	34381	Α	4054	1	705	
4014	34382	Α	4055	1	1833	
4015	34383	Α	4056	1585	4128	
4016	34384	A	4057		1425	MARG/NAITLPV/CGRAVKFT/L EVLRGDSVEKTSRVWSGNERD QELLTEDALDDLIPSFLLTGQQT PAFGRRVSGVIEIADGSRRKA AALTESDYRVLVGELDDEQMA ALSRLGNDYRPTSAYERGQRY ASRLQNEFAGNISALADAENIS HKAHKYFVFEANTGTETGYQG EESLFNKAYYGGGTNFFRKESQ KLQQSAKKRDAELANGALGIIE LNNDYTLKKVMKPLITSNTVTD EIERANVFKMNGKWDFADFGT TIKQDFRLLGQTSVDRLLQLSQ GQAVKGNQLLPVSLVKRKTTL APNTQTASPRALADSLMQLAR QVSRLESGQQSSKQKKAIQTAI RKNKEANAVLARLNSELQQQL KGFADFREPPIKQDFRLLGQTS VDRLLQLSQGQAITELCGAKRV GYFGPTQFYIALKLIAAAQSGLP VRIESIKCGNSYDHDYEFELGTL VLPRSLEGFALSLNCGEHYWL

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4017	34385	A	4058	1461	2496	NKRNHQSVCHAFIRIP\AAPMV DSLIARVGVMARGNAITLPVCG RDVKFTLEVLRGDSVEKTSRV WSGNERDQELLTEDALDDLIPS FLLTGQQTPAFGRRVSGVIEIAD GSRRRKAAALTESEGTPAFGRR VSGVIEFADGSRRRKAAALTES DYRVLVGELDDEQMAALSRLG NDYRPTSAYERGQRYASRLQN EFAGNISALADAENISRKIITRCI NTAKLPKSVVALFSHPGELSAR SGDALQKAFTDKEELLKQQAS NLHEQKKAGVIFEAEEVITLLTS VLKTSSASRTSLSSRHQFAPGA TVLYKGDKMVLNLDRSRVPTE CIEKIEAILKELEKPAP
4018	34386	Α	4059	340	2067	
4019	34387	Α	4060	1	1959	
4020	34388	Α	4061	1	2319	
4021	34389	A	4062	1	1587	
4023	34391	A	4063	1	1757	GYSGSKPDVITLLEQGKEPCVV ARDVTRRQCPAAPMVDSLIAR VGVMARGNAITLPVCGRDVKF TLEVLRGDSVEKTSRVWSGNE RDQELLTEDALDDLIPSFLLTGQ QTPAFGRRVSGVIEIADGSRRR KAAALTESDYRVLVGELDDEQ MAALSRLGNDYRPTSAYERGQ RYASRLQNEFAGNISALADANN ISRKNITRCINTAKLPKSVVALF SHPG/ELSARAASQRQQCGYHK LHDKQRLLRG*KGNICAKLLNE
4023	34391	В	4065	1	1599	
4025	34393	A	4066		682	MKRAPVIPKHTLNTQPVEDTSL STPAAPMVDSLIARVGVMARG NAITLPVCGRDVKFTLEVLRGD SVEKTSRVWSGNERDQELLTE DALDDLIPSFLLTGQQTPAFGR RVSGVIEIADGSRRRKAAALTE SDYRVLVGELDDEQMAALSRL GGATQAFAKENNQK\HTKKRT ASLILHAMICCRSLNSSKTKNT KCLNSINQRLKILSLQKDLMCG TAGRCKTLTEQ
4026	34394	Α	4067	1	2448	
4027	34395	Α	4068	1	2541	
4028	34396	Α	4069	1	828	
4029	34397	Α	4070	1	1899	
4030	34398	В	4071	1	1686	
4031	34399	Α	4072	1	1437	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	•	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4032	34400	Α	4073	1	3417	
4033	34401	A	4074	1	3826	
4033	34401	A	4074	812	3826 2578	FIRDFADFGTTIKQDFRLLGQTS VDRLLQLSQGQAVKGNQLLPV SLVKRKTTLAPNTQTASPRALA DSLMQLARQVSRLESGHSNGN GQVSPIFHQTSSSTIRSCSCHLLT LNFLTLQLNTSDIAVFHSTPKLL LVTSTITHMGLNTSQAQSVPVI NSVAGSLAALQPVQFSQQLHSP HQQPLMQQSPGSHMAQQPFMA AVTQLQNSHKFSHRSHGPGQS NDACSEPTNKKMRRNRFKWGP ASQQILYQAYDRQKNPSKEERE ALVEECNRVWQARRLGAFGKE DVHVSFAARRGAKFRHQTLLG RRSSIPAAPMVDSLIARVGVMA RGNAITLPVCGRDVKFTLEVLR GDSVEKTSRVWSGNERDQELL TEDALDDLIPSFLLTGQQTPAFG RRVSGVIEIADGSRRRKAAALT
						ESDYRVLVGELDDEQMAALSR LGNDYRPTSAYERGQRYASRL QNEFAGNISALADAENISRKIIT RCINTAKLPKSVVALFSHPGELS
						ARSGDALQKAFTDKEELLKQQ ACKL\HEQKKAGVGDNSIDSW KNAGRVFKDSDKFDANDPILK DQTQEWSGSATFTSDGKIRFIL

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SEQ ID		Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	1	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
4035	34403	A	4076	1474	3367	REEGANSECLGRHGFKKMLYV
						KRDEVGKGQIRLETVFEQAIDQ
ŀ				ļ		RFSTDTSLSTPAAPMVDSLIARV
	1					GVMARGNAITLPVCGRDVKFT
				į		LEVLRGDSVEKTSRVWSGNER
	•					DQELLTEDALDDLIPSF\LLTGH
į						KTPAFGQRVSGVIEIADGSRRR
						KAAALTESDYRVLVGELDDEQ
						MAALSRLGNDYRPTSAYERGQ
						RYASRLQNEFAGNISALADAEN
						ISRKIITRCINTAKLPKSVVALFS
						HPGE\LSARSGKCMVPTESAPH
					,	VTVLGCQCGPLGLENGLKEGY
		:				LGRSTLDMEAWQPLQEFYLHN
		ŀ				LITGQMFEIAVTQNNSKINSSSP
						TTEQSWMENDFDELTEVGFRR
		:				SVITNFSELKEHVLTHRKEAKN
						LEKSDGENGTKLENTFQDIIQE
						NFPNLARQVNIQIQEIQKTPQRY
						SSRRATPGHIIVRFTKVEMKEK
						VLRAAREKASLAPENLDNSKIR
						PVVILFHYGESWNLLRADQRLI
						FAKSWPRASRYQQGHQDLFILR
	,					SDLPSQVFIRDKLMERRNRRTG
						RTEKARIWEVTDRTVRTWIGEA
						VAAAAADGVTFSVPVTPHTFR
						HSYAMHMLYAGIPLKVLQSLM
						GHKSISSTEVYTKVFALDVAAR
						HRVQFAMPESDAVAMLKQLS
4036	34404	Α	4077	794	4235	RVSRGRKWFFIALKRMPAMKK
						AMNLFLGLSNVRTVHPEGFTV
						YISTHISFPSLSGYRTGLRSFGLV
						KQKKSPIRMPCVYTNTLCQYR
	1					KPDGSGIVSLKIDWIIERYQLPQ
						SYQRMPDFRRRFLQVCVNEINS
						RTPMRLSYIEKKKGROTTHIDL
						ALKGLRVLLVEGNDPQGTASM
						YHGWVPDLHIHAEDTLLPFYLG
						EKDDVTYAIKPTCWPGLDIIPSC
						LALHRIETELMGKFDEGKLPTD
						PHLMLRLAIETVA
4037	34405	A	4078	1	2574	
4038	34406	A	4079	1	536	
050	121100	14.	1.077	<u> </u>	1	l

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
4039	34407	A	4080	368	1449	LKSTNLITFLRLFVPQMPLAYM
						KFTPSGLVAACAPWLRQRRVV
						QLGQIAFSAP/YLSQMVRQEMY
İ		İ				NRYGESAYEDGYRIYTTITRKV
İ		ŀ				QQAAQQAVRNNVLDYDMRHG
		ŀ				YRGPANVLWKVGESAWDNNK
	ļ					ITDTLKALPTYGPLLPAAVTSA
		ŀ				NPQQATAMLADGSTVALSMEG
İ						VRWARPYRSDTQQGPTPRKVT
						DVLQTGQQIWVRQVGDAWWL
						AQVPEVNSALVSINPQNGAVM
						ALALLNNARPWYLGAQPRDSTI
	<u> </u>					IFCQFGAHPLLDPKTQPVGCRN
						AARKSCAEIRLVPDARANSGKL
						VRRYRKYRRQYHKSKSRHQPL
						RQQQPVRLDWRNVDNQYALT
						TRFLYQSLQRHAQLNVPLFHVL
4040	34408	Λ	4081	1420	1842	
4041	34409	Α	4082	407	1347	GRIRVHIHKDGRADGGSQPGVT
i.						AIQQQLPFAFAFPN*SY*TESAW
		1				AQSIK\GPWWLRDQVDGPAGR
		l				LAALPQR/SLINAVSTRMEGISG
						AFNTANPACST*FLCSLLILPSLF
						STALPNFRLSAMVSDCTISNMV
		l				WST/SAVTDWSCTPLD*ERKHR
						GTARLTTGKGVGMDRDKQVST
						LFLGFCYAHLQWNEDVFIARH
						VHLHIALFLDQRAQTASYLQYH
						IFFARFVFPHRTGVFATVARLK
						HNDNRTIAPCFTRLWTTLRWR
						HLLFEVAFVVILQQRQQRVLHI
						LCIGRIEVHHQTLFKPGDRRKG
						KQLRFYVLL
4042	34410	Α	4083	1	649	MRHGYPARANVLVKVGESAW
						DNNKDYRYAKALPTYGPLLPA
						AVTS/ANPQQATAMLADGSTV
						A/LSMEGVRWARPYRSDT/QQG
						PTPRKVTDVLQTGQ/QIWVRQV
						GDAWWLAQV/PEVNSALVSINP
						QNGAV/MALVGGFDFNQSKFN
						RATQALRQAGAHLPAHSQSGH
						HQQTAR*KSNFCARM*TPDQLS
}					1	W**KNCPFRLSPT*QRQWSLRR
						YRPVSQRTSF

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence		*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4043	[34411	IA	4084	2	551	WRAAGPEPCPTGRQLRP/AGDL
4043	154411		14004	1		ACSAAAGPGAEPFTAPLALAGR
		İ				SKCGAAEPAPTQNSRWPMSPH
		İ				LSLHASPQAEGAGSGL/VPAPR
						AAAG*RAPQMRP/VVGAEAEE
ŧ		İ				APGP*EPCRHPAPHASQHRYGC
						HPS/AAEPCRHPAPHASQHRYG
						CHPSGLNPAGTQHPMPASTGTA
						AIHRG*TLPAPSSP
4044	34412	В	4085	1	1029	Times TETAL SOL
4045	34413	A	4086	1	2157	
4046	34414	Α	4087	1258	1838	TQVVFITSAWGLGEMVVQGAV
						NPDEFYVHKPTLAANRPAIVRR
						TMGSKKIRMVYAPTQEHGKQV
						KIEDVPQEQRDIFSLTNE\EVQE
						LAQQAVQIEKH\YGSPMD/IEW/
						AKDG/HTGNGHVQALRNRCPE
						ARQHRMRIPGRILPRPIGRMAG
						PKTRIEHTSVTVISNRRKIKTEN
						RGHKGYEDRKLHEDLQLRHQS
4047	34415	Α	4088	2806	3540	
4048	34416	В	4089	1	1251	
4049	34417	Α	4090	341	946	GLSSVGQSVNDHLPWT*GLSSV
	}					GQSVNDHLPWT*GLSSVGQSV
					•	NDHLPWT*GLSSVGQSINDHLP
						WA*VLSSVRQSIDDHLPWT*VL
				ļ		SSVRQSIDDHLPWT*GLSSVGQ
						SVDDHLPWT*GLSSVGQSIDDH
						LPWT*VLSSVRQSIDDHLPWT*
						GLSSVGQSVDDHLP*M*GLSSV
					•	RQ*VT*AKVNPKISAVTRNRGS
						VESPHLEGRSLKVQVFIPQVED
1050	24410	ļ	1001	102	T.O. C.	MSWGPPWLWVEGESWT
4050	34418	Α	4091	426	706	VLGGGSEEKAPLWWSGPMVLP
						GAHSMKT*LPHTHVEFGFACLA
						SAGAQDVGMEGPRHTTENSVT
						GSPSHFPPRASQHRRGICRPHAG
4051	34419	 	4092	596	905	RATADF
4031	34419	A	4092	390	905	GLSSVGQSIDDHLPWT*VLSSV
						RQSIDDHLPWT*GLSSVGQSVD
						DHLPWT*GLSSVGQSVDDHLP
						WT*GLSSVGQSVDDHLPWS*G
						LSSVGQSVDDHLPWT*GLSSVG
						QSIDDHLPWA*VLSSVRQSIDD
			:			HLPWT*VLSSVRQSIDD/HSSMD
						VRSV*CRTISR*PSSMDVRAV*C
						RTINR*PSSMDVSAV*CKTINR*
					1	PSSMDVRSV*CRTISR*PS\P*T*
						GLSMSLIPSQLCGLSAVTPFSAV
						TRNRGS\ENHPILKAAASRSKSS
	<u> </u>			<u> </u>	<u> </u>	FPRLKT

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	ı	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4052	34420	Α	4093	3	1194	SLGPRSHSCCRSDYRSGTTVPL
						VLLPVCGVPALLVFALLSPLCV
						VCSALCGGLLPVLRASLFLWCV
						AFLAGLVFVFGFAFFGSLVRGR
						FLVVVPFFLLFALCRLLFLVCW
1						LRSFGACPVSVCVAGFACFAGL
						FLVLVSLLSSGFGFRLSFSCVVG
						SLCLPGFAFRAFCLFFLPCVGPA
						LLAFPGFCGPSSPSLSYGGLFAP
						WSCALLGFFGCLGWSAPGFLSS
						FGLSVRVLSLPCASGLRSLSGC
		[ALVPGLFLPWVFSPRSLRPLVSF
						GCLLCSFVSH\NMDWIKES\AG
						KVIQGNP*WLPVILFFGSVPLTS
	}					KAATAKPLMRMG\RALTVSQL
						T/AVASFAAVYGLFILPT*PTLV
	1	j				GAVQMDDTGTTRIGKLVSNHP
						FFIRVLLGVALTVCFGFVLGSF
4053	34421	С	4094	70	1950	
4054	34422	В	4095	262	4347	
4055	34423	A	4096	2	458	
4056	34424	Α	4097	2	445	QPTERGLCASLKPSRAAIKSQSS
				ļ		KVISFDSMSHIQGTVVQGVGSQ
		ŀ				GLEQQYRSGVAVFRLHSFSHRL
						LSACEFSRCRVQAVSRSIILGSG
						RWQPPSHSSTREWPSGHTVWG
						LQPHISPLHCPSKDSL*GLCLCN
						KLPPENLGFSYVL .

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SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
4057	34425	A	4098		2589	MVWFLKNVNHGTHTNAKKYN
1,057	34423	`	4070		250)	DVENKERTGWKWGSTEYLLCA
		1				RPLRKGNVGLSGDVFLTVFVM
						KTGHSSSLLPSTTTSDSTAQEGY
						ESRGGMLDWKHLLDSPDSTDP
						LGAVSSHNHQDKKAMLKDGEE
						RPFNEPGVFHLLADHQLTQKV
						ASIPGSAVCAYDMLDIASVFTG
		İ				RFKEQKSPDSTWTPVPDERVPK
						PRPGCCAGSSSLERYATSNEFPD
						DTLNFIKTHPLMDEAVPSIFNRP
	1					WFLRTMVRYRLTKIAVDTAAG
						PYQNHTVVFLGSEKGIILKFLAR
	}		,			IGNSGFLNDSLFLEEMSVYNSE
		1				KKWSTAKPVRVTIILNPGQASF
						CITLRETV/C*RRKHIWCPPYRC
	·					TLQ*HFCPCH\CLSGKETLCRVT
						GGMKVKADRDESLPYAAMLA
						AQDMAQRCKELGITALHIKHR
						ATGGNRTKTPGPGA\SRPSSPCP
						LGCLK/WQTLFPRRRLRWPQGG
	İ					RRKRSQLEAQRVIRESYLKGHD
						QLVPVTLLAIAVILAFVMGAVF
						SGITVYCVCDHRRKDVAVVQR
						KEKELTHSRRGSMSSVTKLSGL
						FGDTQSKDPKPEAILTPLMHNG
						KLATPGNTAKMLIKADQHHLD
						LTALPTPESTPTLQQKRKPSRGS
						REWERNQNLINACTKDMPPMG
						SPVIPTDLPLRASPSHIPSVVVLP
						ITQQGYQHEYVDQPKMSEVAQ
						MALEDQAATLEYKTIKEHLSSK
4058	34426	В	4099	1	1299	MALEDQAATLETRIKEHESSK
4059	34427	A	4100	95	502	FPEIPQSCREGAPGPAKPGGPRA
4035	34427	(``	100		302	REPCPNRTAASWGVHCEDGGS
						TVRTGGPL*GRGVHREDGASSP
						QHPPRRGRGLGHLGPRPL*GQG
						DAAAAPGHRGKS/GGKGFLPAL
				ŀ		RVQRGERGRVSRRAVCMWTSL
				ŀ		CASVPS
4060	34428	A	4101	2	653	DSFGSMSVLIKNPRTLFGGKPY
						VCRECGRGFTWKSNLITHQRTH
						SGEKPYVCKDCGRGFTWKSNL
		-	1			FTHQRTHSGLKLYVCKECGQSF
						SLKSNLITHQRAHTGEKPYVCR
						ECGRGFRQHSHLVRHKRTHSG
						EKPYICRECEQGFSQKSHLIRHL
						RTHTGEKPYVCTECGRHFSWK
						SNLKTHQRTHSGVKPYVCLEC
ŀ						GQCFSLKSNLNKHQ\RSHTGEK
L		1	1	<u> </u>	J	- COLOBITOR INTERVENTION

SEQ ID NO:	of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4061	34429	A	4102	8	389	LPASASQSVGTTGVISFENTCNI CHFFFLLLFFSSSSSFFFLPFSFS FRVSLF*IQPLKTTASTVQGRQH SGGYRASPERRADQRAHTGEK P/YVCRECGRGFRQHSHLQLSV YPHSFWTQDKRSNH
4062	34430	А	4103	1	740	EGKRGRPFRIPRPAHPFPHELQS RLCILKPLHHPTPSSCPSS*TPVA PLPSHPCAPH/SARSCPVSDEAA L\P*SMALWARAGLLEAPTTLL PAAPDASSPA/MPPSRKLPPPLP L/CPEPLGPSPAAPSPPAPPGPNA AARPPL/PPSPSAAPGPRRPGA\R PVGPSRGPG/PRNSRSLRAPDVH TAPMRCLPSVRPPLPVLSAL/PD PLPRPPSFVPSLPSP/PSSGPSCPP TSAPPGSPRPGFVRLPCLLFWGS
4063	34431	В	4104	48	272	
4064	34432	Α	4105	2	622	CPLSPLLFNIVLELLARAIRKEK */LKGIQIGEEEVKLSLFGDDLIV YLENPKYSSKKLLELVNEFNKV SGYKIYVHKSVALLYTNSDQAE NQIKNSTPFTTATSSSSSSSSSP QGIFLTKRLKNF*RGKFKTLVK KNQGDPKKGKNPPGPKMGKN NFGKTPFWAKKI*KFHSI\PKKT PPFFFQKLKKTGVKFFWAPKGP KGFLSKK
4065	34433	A	4106	39	1043	QKQPVWQRCREIGTLGYCGWK WTLDIHGRGHRILSGGVEIPGP WTEGFIQGRDVGELQEPGLSGR ESIH*\GKSYEYECSEDGEVFRV RASLTNHQVIHTAEKPYKCTEC GKVFSRNSHLVEHWRIHTGQK PYKCSECDKVFNRNSNLARHQ RIHTGEKPHKCNECGKAFRECS GLTTHLVIHTGEKPYKCNECGK NFRHKFSLTNHQRSHTAEKPYK CNECGKVFSLLSYLARHQIIHST EKPYKCNECGRAFHKRPGLMA HLLIHTGEKPYKCNECDKVFGR KF\NLTNHQRIHTGERPYKCNA CGKVFNQNPHLSRHRKIHAGE NSLRTLQME

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4066	34434	A	4107	3		QQHQQQHFVGQVAIQQQQQQGPGVQTNQALGPKPQGLMPPSSHQDLLVQQVSPRPPQGPQGMVGPAQVGVLQIIQLHGALGPQGLH*QVFMP\QSRVFSSPQLAQQGQGLMGHRLVTAQQQQQQQQQQQQQQQQQQQQQQQQQQQQQQQQQQQQ
			·			GIISYFTQHSWEVKVFTTL
4067	34435	A	4108			MEKNKVVKREAEANSINLSVY EPFKVRKAEDKLKENSDNVLE NRVLDGKLSSEKNDTCLPGTAP SKTKSSSKLSSCSSAIMALSAKK AASDSCKEPVANSRESSPLPKE VNDSQARAPLQSTVMTNAVSP AELTPKQVTIKPVATAFLPVSA VNEMKTAGSRVINLKLANNTT VKATVISAASVQSASSAIIKAAN AIQQQTVVVPAPSRANAKLVPK TVHLANINLLPQGAQATSELRQ VLTKAQQQIKQAIINAAASQPP KKVSRVQVVSSLQSSVVEAFN KVLSSVNPVPVYIPNLSPPTNAG ITLPTRGYKCLECGDSFAVEKS LTQHYDRQSMRIEVTCNHGTK NLIFYNKCSLLSHARGHKEKGV AADTRGQKTCTICQMLLPNQCS YASHQRIHQHKSLYTCPECGAI CRSVHFQTHVTKNCLHYMRRV GFRCVHCNVVYSDVAALQSHI QGSHCEVFYKCPICPMACKSAP STHSHTYTQHPGIKIGEPEIIYKC SMCDTVFTLQTLLYRHFDQHIE NQKLSVFKCPDCYLLYAQKQL MMDHIKSMHGTLKSIEGPPNLG INLPLSIKPATQNSANQNKEDT KSMNGKEKLEKKSPSPVKKSV ETKKVASPGWTCWECDRLFIQ RDVYISHVRKEQGKQMKKHPC RHLCQHNRIKHKGIRKVYACSH CPDSRRTFTKRLMLEKHVQLM
4068	34436	В	4109	1	411	
4069	34437	С	4110	54	146	
4070	34438	Α	4111	1	1937	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	1	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4071	34439	I A	4112		1830	MCIEVTCNHCTKNLIVYNKCNL
4071	34439	A	4112	1	1830	
ļ						LSQARGHKEKGVVMQCSYSIL
						KPVSAGHIIVSPSSNSSSSSSTLQ
	}					SPVGTGIHTVTKIQSGITGTVISA PSSTPSTTAMPLDEDPSKLCRH
						NLKCLKCNEIFQDKRSLATHFQ
						QAADMSGQKTCTICQMLLPNQ/
						CQRIHQHKSPYTCPECRAICRK
						KRT\QIHEWERETGKEISISFEKK
						SMETKKVASPGWTCWECDHLF
		•				MQRDVYISHLRKEHGKQMKK
						HPCRQCDKPFSSSHRLCWHNRI
				l		KHKGIRKVYACSHCPDSTGTFT
						KGLMLEKHV\H*CMASRTLT*K
						K*QTPPMRRKQK*K*TSRSAVP
						SG\VERTGSGVQASQRSNNSTT
						EKAENQCF*GSQAPLCCTQVKG
						TSASAQAKWGWRR*PTGEQTQ
						PRGRISQW\VMSDRKCKVCAKT
						FETKAALNTHMQTHGHAEGCL
						KQPCRSLLSQPRIKTEARNLIRN
						ADFLNSILRNGEGYSKEKKNGT
						GFLGRSARLALGAQGGKSWRF
						LFWVLLPNVLVLRVGMHDVN
				}		HRLINAAGCVSQLAVTLSTEPH
				}		GISSAISRVPRHCHPSGENSMAT
		ŀ				SLNVNRSISRLAAGSGVLAMDL
						PIPAGHRAIETGLLGTEDTEQ
4072	34440	С	4113	217	510	
4073	34441	A	4114	210	281	
4074	34442	A	4115	1	675	
4075	34443	C	4116	126	434	
4076	34444	A	4117	804	2061	WERREAGGEDEGINIHEP*VEE
					!	EMKKHESNNVGLLENLTNGVT
						AGNGDNGLIPQRKSRTPENQQF
						PDNESEEYHSLGDKSKTSFQNS
						NNNNNKQQEQQQQNPTFSNTR
						KLTKLYKAPIPPSIILSGCPNIND
						SNWQEIEHGMQTAGLPTRPLSH
						GLQQKGAAFRCLGCKCSEPFTG
						SLILQKAKTNTQKWQATYPKS
]		QNEQLVPSVGKSYRCSTPAQP
						MKTAVGHKPCKATGAELPKAL
				1		GAQPLHPCALDVGQGFKKGNF
						GAVGLNGLLGLEFHGVSGVLL
						VGPGDGGLISEGVVREDLMCG
						VWSAGTWSVGTAERCLEKPGA
						LHVIEGPLDSWDGPVMPNGPV
						KSRQSSCLDGPGRCCSEILTGQS
						HGNKKPARASSKSSQSINDRPL
L	<u> </u>			<u> </u>		AVLTNQYQCEQLASERQPSSNS

SEQ ID	SEQ ID NO:		SEQ ID NO:		l .	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4077	34445	A	4118	1	357	GKLMLPGSTRKPQVPVDRKMG
						HEGVFVWQAGLRARPGPSLPFS
						HGLTLHLHWPLALPV/GATSSP
						CEAGDLGV\PLAAGTWCPWEA
			}			*RQEEAGWQAPGRAGPARVG
						WGGTGLTSAEVIITI
4078	34446	Α	4119	1	771	MLISHKQLSPQLLSLTLPLSEGK
						AGWAGECPNPPSKDSPVQGIG/
						GPPP/GYQKCR*DTASMLTMAP
l						CHGPVCPPHPGWRPRSGSRVIL
						PAPPGHHPWP\GPRARNGLGTF
						QGCSTGWQVQETCFGPGGWLG
						DRHLVGPAATGARCLPAARGP/
						DGALHPAVPTGKLKGQP\GPGA
		ŀ				RHQRTYD*LPRPCGAAGLGSPP
						A*HPISEETENQWGLHGPPQPA
		l		ŀ		WARPDHGCQ/APTLSPSLKRKP
L						GRVTAGGPMPGCFSTSSVPTT
4079	34447	Α	4120	1	402	MLISHKQLSPQLLSLTLPLSEGK
						AGWAGECPNPPSKDSPVQGIGD
						LHQLPKMQIRYSIHADDGSVPR
						AAHKRGLRKRTLKTISLPRQES
						AFPFHGQGGDPGVVPGSSFLHP
ļ						LGTTPGQGPRARNGLG\PSRAA
						PPAGRGVAGGSPSGG\PAATGA
						RCLPAARGPVGPYTQQSQRAS*
						KGSLGPGARHQRTYD*LPRPCG
	•				ŀ	AAGLGSPPA*HPISEETENQWG
						LHGPPQPAWARPDHGRQPLRC
						HPP*RGSQGGPQKAAVSQQIPR
						AGQEGTH*DPTEWGPPDGDQG
		1				GPRESCRGLQGGRGQCLCDWS
						PNTSEI*YPHA*NGD*KAG\PPM
				1		DTKSQLQVSTPKSPASHGEDVA
						RLEEPEASGD/RSVP\GLPGASLI
						PIWRPPFSRISVRTFLPSPWNLL
1000	24440		4101	111	010	RDCGFLGTSLASSSGRVTA
4080	34448	C	4121	111	218	WWWW CAMBECH BODI BOOFA
4081	34449	A	4122	2	453	WWPVLSVPPECRLPGRLPSG*V
			1			RGPAPWWPEPASQDKSQLSSR
						GFPGKVSLGKGMAFSPLQTAP*
					•	KWLGLSPPLSSTENTASRGHTS
	1					PSSRNGFDSQPRDSRTGRECQA
				1		TQLPAQHSHAEVLHFGGAMSG
L	<u> </u>	<u> </u>	<u> </u>	I	<u> </u>	QLSLVGPQDSKRTARLTDSQ

	-	l .	-	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/340,217	sequence	or pepade sequence	deterion, (-possible adelegade insertion)
				'		
4082	34450	Α	4123	1146	1775	KGWLEGAPEA*ERYPG\PPCVV
						CSLSDAPWVGHPAPSLGV*EP
						NPP*SRGESQGPTSSICRQSSGD
						LG*KLESPIHTVIPRHTSQARQG
						HTAPFPPFQVPS*LLV*LKAVSL
		ŀ				APAEAP*PASGLHAPWPVAPRV
						TCAI/PAKGTTVPAGPAELRPVS
						PPPILLP/PDSRSSAFPSRKGPASP
						YETCSPPTS*/EVPS*TYKSMGP
						GIRLPALPASPRVPSEGPGLSEH
						PEGPPALPPAIPFS\SPSWFKQCS
						FSIRPGWLLHGAPQGKGWPQA
						SWVGKTG*PEKRGSPRGPEHSA
						LNLRVALPGVAV*EGPACVGW
		İ				GGPPQPPGAICEATAPPSI/VPPL
						SLPAPFPGTLP/PPTPAASP/PPAL
						PPLLRRGRPRPCAALALPALSSL
						FS/PPVFSLLSLQLPADRVRQVH
						PVLRAPGPPFRPPKQIPPSSSGDL
			:			PFPSLPGR/PVL*LEKWLPPAPK
						ASPPSSVNLILLVVVKLNTFRCG
						PLVKNLVPPSVVCPCPCSYKYL
						*ILIYIHTLHMGQPPSPSSAGNQ
						SLCYPCGGLVAQPTKRTLVPPTI
						QLQSVPPPV/KPPCHARPVDSQP
						PPSLPPPTKHGGAVQAAVWPDS
	:					FYPVLLSLG
4083	34451	Α	4124	146	1701	TFLGYLETAHGPSAQQCPTGLF
						AFRSLGRGLLLTSLPKQPARSPP
						REDVPRSTTQEMTRPRHPPRKP
						AQPGLGARRRGAPV/RGLSKSR
						ELNSGNTSDSGNSFT\PPHPRTR
						GPCWRISPPPAGAESQGDAMLL
						ARMCQMPSLGLMSRTFPHSST
	:					GKARGFQSPCLECAEVKKSSLV
	İ					PSTARSSPMKGCSRSSSYASTRS
						SSHSSQSPNPRASPRVRTIITCIL
						*TRKRPRETKSSAKVT\HYYSSK
						SGKRSPPSRSSRSRRSPSYSRYS
						PSSPNSPADIPQNSHPQPSASTD
						RPHIQSPQFLPTHQGLRNIHVLT
						PAAPALL*CPPANADTPAQAQP
						PPLRY*QPSQTLTAAPSSSLRSP
						LRQRADPIP*PSGGAGSQIQ\WK
		1		1		DSQQRERERARRRRRSYSPMR
		1				KRRRDSPSHLEARRITSARKRPI
						PYYRPSPSSSGSLSSTSSWYSSSS
						SRSASRSYSRSRSRSRRRSRT
						RTSSSSSSRSPSPGSRSRSRSRSR
			1			SRSRSRSQSRSYSSADSYSSTRR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
	<u> </u>	<u></u>				
4084	34452	Α	4125	1	1068	MLLLELNAPEHVLETINFQTLT
						AFCNTFHILRPTKAPGFVYAWL
						ELISHRIFIARMLAHTPQQKGW
						PMYAQLLIDLFKYLAPFLRNVE
						LTKPMQILYKGTLRVLLVLLHD
						FPEFLCDYHYGFCDVIPPNCIQL
						RNLILSAFPRNMRLPDPFTPNLK
						VDMLSEINIAPRILTNFTGVMPP
}		1				QFKKDLDSYLKTRSPVTFLSDL
		l				RSN\LQVSNEPGNRYNLQLINA
						LVLYVGTQAIAHIHNKGSTPSM
		1				STITHSAHMDIFQNLAVDLDTE
		1				GRYLFLNAIANQ\LRYPNSHTH
						YFSCTML\YLFGRRANSGRPFQ\
						EQITRVLLERLIVNRPHPWGLLI
		İ				TFIELIKNPAFKFWNHEFVHCAP
4085	34453	Α	4126	1	984	MQANLEMAGNGVTSMGMEPL
		ł				AIPHIYCCSEGTCNFSNTENHCL
						RAALSMLLNGTPFAFVIDLAAL
						ASRREYLKLDKWLTDKIREHGP
						SVHGLFPSRVLSPALGPGAFPG
						RHNCGSCVAPQSGLPGVHPVEL
						PWSISKLFRLRSPANFSDVLGSR
						SKVLLLMCTLKYCGMQLGADA
						TRVDMLTFLPTLGFIRNNDYTD
						DTKASELTELSHNLHAYDSVTG
						VPGDETECSKTVSTWAYTAESL
		ŀ				QGYMAAKLLGRNLTVPSRYLF
		İ				LNAIANQLRYPNSHTHYFSCTM
						LYLFAEANTEAIQEQIT/RLVRE
		ŀ				RI*S*ANAYWHSEKFYQFTCEL
4086	34454	С	4127	1	399	
4087	34455	Α	4128	I	868	MANVCNPSTLGGRGGRITRRPE
						DPGSPVYSVPPASYHPKPWLGA
		ŀ				QPATVVTPGVNVTLRCRAPQP
,						AWRFGLFKPGEIAPLLFRDVSS
						ELAEFFLEEVTPAQGGIYRCCY
						RRPDWGPGVWSQPSDVLELLV
		1				TEELPRPSLVALPGPVVGPGAN
						VSLRCAGRLRNMSFVLYREGV
		1				AAPLQYRHSAQPWADFTLLGA
ŀ		1				RAPGTYSCYYHTPSAPYVLSQR
						SEVLVISWE\TLAPPTTPGGT*S
						AWGWPGWSSSPWARWSLLTG
						AVRTALLLFPQVPHRATTPWVT
-						SYDWVWLP
4088	34456	Α	4129	1	270	
4089	34457	В	4130	39	919	

SEQ ID	SEQ ID NO:			•		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN 109/540,217	location of first	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/340,217	codon for peptide sequence	of peptide sequence	detection, \-possible nucleonae insertion)
4090	34458	Α	4131	3	466	GRALCPPRLLAAGRVLGPGRRS
						PG\PGPGVP/GG*R*GGAEPRPG
'						APRGRVLPPSAGSGQFSAATPA
						QNGLPALRGPGSRPGIRSKAVR
						PVPLGRVGVYFRDALRASGQS
						GRKLCCIGNTLSPTSFSVGKEVP
						RKHETNQKHEKGILCMEAVKP
4091	34459	Α	4132	1	1647	MWRWLYAGARMTVRDKQPLE
						QMLAGCTHASLVPTQLWRLLV
						NRSSVSLKAVLLGGAAIPVELT
						EQARDMGIRCFCGYGLTEFAST
						VCAKEADGLADVGSPLPGREV
						KIVNNEVWLRAASMAEGYWR
						NGQLVSLVNDEGWYATRDRRE
						MHNGKLTIVGRLDNLFFSGGEG
						IQPEEVERVIAAHPAVLQVFIVP
						VADKEFCHRPVAVMEYDHESV
						DLSEWVKDKLARFQHLVRWLT
						LPAEPKNGGIKFHVSAKRVGAL
						TTRMEAAQQHADDKIRQMINS
						EQRLSEQFENLANRIFEHSNRR
						VDEQNRQSLNSLLSPLREQLDG
						FRRQFRTASLMKVAGWDYLM
						NSLYNANSSALVNRVRYKWIA
						AFEGGFTGIVATLDTGRPGPVM
						AFRVDMDALDLSEEQDVSHRP
						YRDGFASCNAGMMHACGHDG
						HTAIGLGLAHTLKQFESGLHGV
						IKLIFQPAEE\VRVARGRWSMQ
						VS*MMLIILLPCTLALAYLRALL
						CAAVIILWQPPNLTRTSPVPPLT
	ļ					QAQNQKTVTMPCWRHTSHSCT
						ACNRPAQRRSFQS
4092	34460	Α	4133	864	1128	TGRSTIRRQRREPRRKAATLRF
		l				DRNGCRARCTPP\GRKEQRYQQ
						TADGDKGAEFYRRPEGVEIVA
						VMEQRDEVIQADKLAGETKRI
İ						DAL
4093	34461	A	4134	618	1102	HSNAAPTARSSFVQNTPSSCGY
						SRRAWRISSIPEETDRTSYRNIQ
						CGNHPPVPARLPAGQTGDGDIP
						PRPRWQSPQTAPRKPPDLP/LIR
1						KNKIPMFRSATTQPGLTGTIAK
						ASSSEVKAIIGARVKMTRSENF
					1	GIQSSLKNILIMSATSWSEPPQP
						TRLGP

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4094	34462	A	4135	2400	3201	VGGGRNRSPGVACWVEDGNG
						DAVRDGW\ALDADWQASQAM
						RRKQQPE\YWQHDAATADADR
1		ŀ				QMNMPV\LH\RALVAGRAQEM
		i				NGAQGKTLLKGW\DEASARRG
						T\KTED\FCEEDTRDDAMIAWM PR*PGVLPRADAWANVLNHGV
1					•	GWKKAQLK\STWM\PQVQQ*W RGD\KAANSNAAVAHTVVLLN
		į				SGGDATQ/TAAFPISSPLSV/EEV
			1			CVTMVIA*FWMWDSGSGGVVL
						CSSSGRSLTSWSATKVRG/SGH
						KGRWCSRQGVTCQVRHGGHV
						APH
4095	34463	A	4136	118	1008	Atti
4096	34464	A	4137	3	1140	KHTYMLSILKPVRTSALPPPAP
1,020	34101	` `	1137	[1	AQTLCTQLSRVSSRLL*DHPSR
ŀ						WGLR/PSTGMSQARACSPGSLG
	-					WMQRSSFTPGAGRVRHIPN\SA
						GSTRRPACGSRSAAAPVRPCRR
					 	TR*G/RSSVVERFMTALSLACR
						ALPGP*AAPGPSITRRFTISAELK
	·					DTRL/PREHVVPLVCTHAIAVPD
						RGAAVRPTRRRDAAAPPSPLVG
						DVTLQCPSQ*RGSNAPDQVRLP
						CVG*RPRSSLQRSGLSVFSADGS
						TSGPEPASGRKDAGWPARVLRF
						GTLSRGAPEAGADWGPYSPGSP
						GAAASGAPWLGQPQALQGAG
						GQLVGSENGERGTGTKPRVSVS
			:			VAYGEIALPADTSWSSRAGAA
						VLLGLSRSTGGEGPGNMGHGG
						QSQMLTLEVL
4097	34465	Α	4138	10	585	PLEMELNLISIEVWGERLGISTG
						TEKMPTLKHRTWPVECSKASSL
1						EGDLRSL/S*LEVISAFSADPASA
1						DDSPGCWKKKDDCSMVHLHR
						QEWQQQQCCQ*K*RKQPPGER
						RNKCGSHPVCGTVLWQP*QTH\
1						QISSCPTVGCPPSHSSFSILDGAN
						AGQEKQSTTEPEPALFLLPPSRG
						AFGPFGLLSDLRRQL
4098	34466	Α	4139	1	474	
4099	34467	Α	4140	458	612	ASCMASVDISVLTCMWRCTIEQ
			1			SSSFLCLLTPLWE*SWCHVTRIC
	<u> </u>		<u> </u>	<u> </u>	<u> </u>	PFISLG

SEQ ID NO:	of peptide	Met hod	SEQ ID NO: in USSN	location of first	codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4100	34468	Α	4141	11	829	MGRPDLILSSILGLRWLLLPQSS
						RAGRRGAIREASDAEQVTFSGG
	}					TVPARTSSGREWRSLLGPDMET
	1					TSLFCIDTKTTFLFVYRYGDHIP
						LVPEQSGLEPLLIWADPGLFHV
						RLFHLLTIDDNFCGLDMNAPLG
						VSDMVRGIPVFTEDRDRMTSVI
	l					AYVYKNHSLAFVGTKSGKLKK
						/VRSSAAP*VWDQDTPRSLGQQ
						GAQRQP/ILRQCVYTFFQLN*RC
		ļ				PHSTARRELRTGGFIPTRSHQD
		1				GLRSAARCTQGTTQ*ASWSCV
						HVCASVRAMCSYC
4101	34469	Α	4142	5	237	NFGAMTRIR\DLPWEINPLSSCS
	155					SLCEKDPPTTSSPQTN*PKEHHT
						NFQSETGDEFYPWTQNFSTGHG
						LGKTVFPWCL
4102	34470	A	4143	1125	1190	2011, 111 1102
4103	34471	Α	4144	306	573	RNFGAMTRIR\DLPWEINPLSSC
						SLLREKDPPTTSGPQTNQPKKH
						LTNFKSGKRPLLTLFSNLSHCPS
						TTFFFPFFNLSLLLISIPFIFW
4104	34472	A	4145	1	329	ASHSWQTLQHSGRYSRSSG/SA
						GSPRDCAARAPTISPGCAMAWL
		:				NLDSISPSSQSKASPLSQLTCPET
	ļ					SYTGCP*SAPHSPPPPWCPQERC
İ						ACKGHCLHHRDGCCGYGYN
4105	34473	A	4146	2	336	SILTRKCKYGMEIPT/NIPGLGA
						AGPTGMFFGSAPSPMGGISPAM
						TPWNQGATPAYGAWSPSVGSG
						MTPGAAGFSPPKA/PTYSPTSPG
						YSPTSPTYSLTSPAISPDDSDEE
4106	34474	В	4147	1	1260	
4107	34475	A	4148	150	335	SFQQSAPW*ASGQSCASDPAPP
			1.2.1.2			ATARGREGPHQSQAFHSRHSPIP
						DPLPPPCSGGWGHSRW
4108	34476	В	4149	1	3267	

SEQ ID	-	1	SEQ ID NO:	4	1	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		07/340,217	sequence	or peptial sequence	deterion, (-possible nucleonde insertion)
4109	34477	Α	4150	1528	2973	GKQIFSLIIPGHINCIRTHSTHPD
						DEDSGPYKHISPGDTKTVINNW
[LLIEGHTIGIGDSIADSKTYQDIQ
						NTIKKAKQDVIEVIEKAHNNEL
						EPTPGNTLRQTFENQVIAVVGQ
		l				QNVEGKRIP*LPPPL*WGAGRR
			:			EGVG*CGRFSEHSRSSAAGYRG
		ŀ	;			GRPCC\PSWEDDP/GAPPDPPAS
		l				AQIPAPTRSGCVRCSARPSPGPP
						AECAAGPSHWYNQSGTSSQKP
						CWKA*AYPGSGTQRSRGSQAR
						SHP**SEAESGA*SQMESACPRR
						SAVQRQQ*PDSQTSAAECSLG
						WAPAHCCVPSR*PLLRPAWPS*
						*CSECPGKS*NQQWSPQCQ*YD
						PNPQSTVVAEDQEWVNVYYE
						MPDFDVARISPWLLRVELDRK
						HMTDRKLTMEQIAEKINAGFG
						DDLNCIFNDDNAEKLVLRIRIM
	<u> </u>					NSDENKMQEVMGVLEVSVSHV
4110	34478	Α	4151	459	940	HLPGGGVPGREGGSPDQHVAP
						GAVSGGAGGGSTRGRGSRRRR
						PGRPRPGPRQPRRGALPGGEHG
						LRASARCAARAQQRDPG/TPSC
						SSWACPTPRRPWAPAASSRLRR
						PPRGPACATPPPCRPPARRTCTG
						RCPPSCCLCGSPTTWRRPPPTG
						GALESPKRR
4111	34479	A	4152	264	1386	SSRCQPVCESGHPGYGQSPA/YT
						TAGRTESGGTGST/GDNHPLWP
						CI/GGAPCPAQNTPHLRCV*RSH
						ALALDSAGSSSPESPH*RASIPH
						TTLGQKRRSWAGTAHS\PMAPC
						AAASISTST*LSHHHSPAAQSVC
ļ						PSSHTTPSFCPIQKFHCFR/SPQR
						NTS*VVLCPGG*LRVG*WPSSG
						HDRSWYHTREPSVGN*HRSHQ
						RR*RGTAPAPGPSARLQCPARG
1						SRSSHSAPASSSRRPFPGSTPAG
						LGFPSARFPVGKPVVPAALMNR
						PTRGERRFAYWAPGWFFFSPVR
						RATADCPSP/SWP*ESCSKRSTL
						VCPSRRKSCLMVVPKSAKSPVL
					·	AK/YGPVGHDHGTEHDVVLGE
						VQGKRPVAPTMGTPKHKAVHP
	5.1105		11.50			RAPH
4112	34480	В	4153	52	363	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4113	34481	A	4154	321	802	HLPGGGVPGREGGSPDQHVAP
				<u> </u>		GAVSGGAGGGSTRGRGSRRRR
						PGRPRPGPRQPRRGALPGGEHG
						LRASARCAARAQQRDPG/TPSC
		1				SSWACPTPRRPWAPAASSRLRR
:						PPRGPACATPPPCRPPARRTCTG
						RCPPSCCLCGSPTTWRRPPPTG
:		1				GALESPKRR
4114	34482	A	4155	15	263	CGRFSEHSRSSAAGYRGGRPCC
						\PSWEDDP/GAPPDPPASAQIPAP
		ŀ				TRSGCVRCSARPSPGPPAECAA
						GPSHWYNQSGTWKCKG
4115	34483	Α	4156	3	518	SPSVGSGMTPGAAGFSPSAASD
]		ASGFSPGYSPAWSPKPGVPGVP
						QVPSKPLKSLHPGGVVRHLSGQ
						VCFLHSSG*VCPLLISFVGCFPT
						GGAMSPSYSPTSPA\YEPRSPGG
						YTPQSPSYSPTSPSYSP
						TSPNYSPTSPSYSPTSPSYSPTSG
-						SYSPTSPSYSPT
4116	34484	В	4157	620	6763	
4117	34485	С	4158	430	870	
4118	34486	Α	4159	1	3039	MDSETRRTAKVRLLMTVLRDQ
		:				DRVSGVQAHPEQFQQAICPLCG
						VSLTRSGTTFGSPSEIYSPLGESR
						ASSGLPRRDGRLIGEEPPEKKFS
						RSPKGD/LSSGGQRIDYRVCVPT
						KFNL*VLSF*PRGQGAGGQSPG
						FSVRRLLVLVWSSGTFV*NGK*
						QKLL*TLCECVHD*GVQGPASG
						SPVCSSTAKATEFEKDPSGPFSS
		İ				SSLPLTPYISFSRVTASSASPGLG
						SALTPQTLKRKGRI*AICL*VVE
						TPKVFR
4119	34487	Α	4160	[1	772	MVARAFLWSQVIRRLGRKGGL
						SQGDRGCNTALAEGRLDPDLT
					:	RGHPALCLPRRPAPRPAEVRRE
				:		GEAEQPEAGQPPGAAPRRARD
						NGAAAAAAGGRLLQSVRPAVV
						CPHPGPQASYGLRYIAKVLKNS
						IHEKFPDATEDELLKIVGNLLY
						YRYMNPAIVAPDGFDIID\MTA
						GGQNNSDQRKNLRSTAKVLQH
						AASNKLFEGENEHLSSMNNYLS
						ETYQEFRFKNVTFDIIATEDVGI
	<u> </u>					FDVRSKFLGVEMEKVQLNIQ
4120	34488	Α	4161	174	444	YHRHDSWRNTRR*\IKLDGKGE
						PKGAEESEATSKYTAAKLHEK
						GVLLDIDDLQTNQNAVNDFSV
						GPQDEVIVEDITNCYLCEIFKRY
	<u></u>					EWVT

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	Į.	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
4121	34489	Α	4162	379	520	GTSHRGASQRRCCPPLSKTGPK
						TPCGKGAPSAP*QGGLDVQGEP
		1				GGK\GDSSGECVGGNVAGLHK
ĺ						GGGTRADSQVPGGMRGGRMS
						Y/GG/HLTAEGPMGRSGPR/GAV
						PSLYPPGFSRGSCSRQYSGAHM
1						PILTGHVGWSESSLDPPRAGQD
						RFLGTARP*GTSHRGASQRRCC
1						PPLSKTGPKTPCGKGAPSAPFIP
	•					AGPTFDHKALM
4122	34490	Α	4163	455	798	
4123	34491	Α	4164	32	2109	WIGGCPGSQPDATAIMGWTLA
						PHSSRCHRCCHYRCHCRCCLCP
						AEMTVGRPEGAPGGAEGSRQIF
		1				PPESFADTEAGEELSGDGLVLP
						RASKLDEVLSPQEEIDPTSDSTG
						SIYHTLLDLAQKGRWLSVWSLS
						FSLTQRVMKTSLKMRRTWRVS
						SKTRTGGWCRSSARRL*GVAPQ
						GAA/DSLNNLPSNIPRPQTQPPS
						GSRPPSQHRSVSSWASSITVPRP
	1					FRMTLREARKKAEWLGSPASF
						EQERQRAQRQGEEEAECHRQF
						RAQPVPAHVYLPLYQEIMERSE
İ		1		i i		ARRQAGIQKRKELLLSSLKPFSF
					}	LEKEEQLKEAARQRDLAATAE
		1				AKISKQKATRRIPKSILEPALGD
						KLQEAELFRKIRIQMRALDMLQ
						MASSPIASSSNRANPQPRTATRT
						QQEKLGFLHTNFRFQPRVNPVV
						PDYEGLYKAFQRRAAKRRETQ
		ł		İ		EATRNKPFLLRTANLRHPQRPC
						DAATTGRRQDSPQPPATPLPRS
						RSLSGLASLSANTLPVHITDATR
						KRESAVRSALEKKNKADESIQ
						WLEIHKKKSQAMSKSVTLRAK
						AMDPHKSLEEVFKAKLKENRN
						NDRKRAKEYKKELEEMKQRIQ
1						TRPYLFEQVAKDLAKKEAEQW
						YLDTLKQAG\RRKTL*ETRVKA
						PGLFKRKRPKSRIFPGSKKLQNS
						ASEIQSR/RLEGSLEQPASPRKV
						LEELSHQSPENLVSLA
4124	34492	Α	4165	251	637	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
4125	34493	Α	4166	1	1344	PGRTTRSMAEDVFLSAPIPRGC
						ADGRDADPTEEHMAQTERNDE
	,					EQFECQELL*CHVQVGAPEEEE
						EEEEDAVLVAEAEAVAAGWM
						LNFLCLSLCRAFREGRSEDFRRT
						RNSAEAIIHGLCSLTACQLRTIYI
						CQFLTRIAAGKTLDAQFENDER
						ITPLESALMIWGSIEKEHDKLHE
						EIQNLIKIQAIAVCMENGNFKG
İ						AEEVFERIFGDPNS\HMPFKSKL
						LMIISQKDTFHSFF\QHFSYNHM
						MEKIKSYV\NYVLSEKSSTFLM
						ľ
						KAAAKVVESKRTRTITSQDKPS
1						GNDVEMETEANLGYKKKC*LT NSLR*LNPQRVQYPY*GSHKNL
-						FLSKLQHGTQQQDLNKKERRV
						GTPQSTKKKKESRRATESRIPVS
						KSQPVTPEKHRARKRQAWLWE
						-
						EDKNLRSGVRKYGEGNWSKIL LHYKFNNRTSVMLKDRWRTM
						KKLKLISSDSED
4126	34494	A	4167	1	1345	IPGSTISCLKGQYPSEPFNMAED
4120	134474	A	4107	1	1343	VSSAAPSPRGCADGRDADPTEE
						QMAETERNDEEQFERQELLEC
						QVQVGAPEEEEEEEEDAGLV
					·	AEAEAVAAGWMLDFLCLSLCR
						AFROGRSEDFRRTRNSAEAIIHG
						LSSLTACQLRTIYICQFLTRIAA
ŀ						GKTLDAQFENDERITPLESALMI
						WGSIEKEHDKLHEEIQNLIKIQA
						IAVCMENGNFKEAEEVFERIFG
						DPNSHMPFKSKLLMIISQKDTF
						HSFFQHFS\YNHMMEKIKSYVN
						YVLSEKSSTFLMKAAAKVVES
						KRTRTITSQDKPSGNDVEMETE
						ANLDTRKRSHKNLFLSKLQHG
						TQQQDLNKKERRVGTLQSTKK
						KKESRRATESRIPVSKSQPVTPE
						KHRARKRQAWLWEEDKNLRS
1						GVRKYGEGNWSKILLHYKF\NN
	1					R\TSVM\LKARWRTMKKLKLIS
4107	24405		41.60	2	270	SDSEDWIVFVKL
4127	34495	Α	4168	3	378	LTSGSRADQGEGQEEGAEGGR
						ASSSSSSSPRGPQHHPHLHGDP
						AEHRPGHPLCSPPDLTVAYQ/M
					1	PEVPAEDM/SDPSFCSARQGGQ
1						RGLDSGPGAPWSSSHSPHSRFQ
L				—		EASHGACAGWRWCRQEEL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	I .	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
		·		sequence		
4128	34496	Α	4169	1	1044	SGQEVNKEDGQADVQQDHHA
				ŀ		DEDGVGDLARQPHLFHLHLRL
			-			CRQCGLLPRLLLGPRLGQPSSLS
						LAGLGPGLGFGDDGLGLQLGG
						GLACRLGASERGGLQRGGGRG
						RGRGLGPG/GPRARAGPQRVGA
						RAAWCAQHSCGSPGKAPPPAP
]						A/TGAGGACRASMAMRSVAGR
		İ				AGLRRPAPSDGVTDRLPSPLGS
				Ì		PFQAP/EAQQAVLGHPGPGPLG
						LRGRPGR*RGATLGPRGLT/PRA
						AAGSRGAAVGGPLRRRPGRGA
						PAGSPSSPGSPAAAGASDIPDLA
						GRSPEPAPWPKCEQCWTPGWQ
						PGRPVPLPQLWPWRGLSISGSM
						PLGEGLEDGSDPMTPSCCLPGT
4129	34497	Α	4170	1	732	SLTQAGTVSLGLDAEGQEVFVP
						FSAVLPMVAPNDLVFDGWDISS
İ						LNLAEAMRRAKVLDWGLQEQ
						LWPHMEALRPRPSVYIPEFIAA
						NQSARADNLIPGSRAQQLEQIR
						RDIRDFRSSAGLDKVIVLWTAN
	-					TERFCEVIPGLNDTAENLLRTIE
						LGLEVSPSTLF\AVASILGGLCLS
						FNGSPQNTLVPGALELAWQHR
						VFVGGDDFKSGQTKVKSVLVD
						FLIGFRLQRP/VSIVSYNHLGNN
4130	34498	Α	4171	1	908	MEKAPPQTQHEGLKSKEHLPE
						QTDEGKTEYRRVPSLRAVVLFR
						QRSCIENILRACVGLPPQNHML
						LEHKMERPGPSLKRVGPVAAT
						YPMLNKKGPLVWEVSPATLFA
						VASILEGCAFLNGSPQNTLVPG
						ALELAWQHRVFVGGDDFKSGQ
						TKVKSVLVDFLIGSGLKTMSIV
						SYNHLG\NNDGENLSAPLQFRS
						KEV\SRSNVVDDMVA\SNP\ML
						YTPGEEPDHCRMGRNLPE*GSS
4131	34499	Α	4172	85	529	ECGARPGSSTRPPARLSPRLFCS
						AIRAALKTRPAPALACTWRTG*
						RASLPTTRCAGSGLGTCTAEGS
						EGCSHPGPLTGTG/RQEACPGT
						APAGSPSCLHPRGRPRPCPPGTL
						APRMSCPWPRSPPLTRYLPSGE
L						NLQSKLESLNTSEKF
4132	34500	С	4173	215	324	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	1	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			:	sequence		
4133	34501	Α	4174	2	505	YKCEVCDKVFNQN*FLVCHNR
					,	CHTGKKPYSCYECGKTFSQTSS
						FTCYRRLHTGGKPYKCSEHNK
						TFG*NSALVIHKAIHTGENPCKC
			•			NECGKVFNQKAHLARHHRLHT
						REKPYKCEECEKVFSRKSHLER/
						HKLKRGGVALV/C*ECPTVYQN
						TSCLRSLSCSYPMSLNG
4134	34502	Α	4175	1	6192	
4135	34503	A	4176	2	3389	
4136	34504	Α	4177	3	875	GEEAALSLCMHSTDDATRLGA
						RDTEPLWHVPAQ/ARLSAIAGS
	1					SGNKHPSR/QDAAGKDSPNRHS
						K\GSKPSAGSLRLSSREGEDRTA
						WTGPRGAVEQEVTGPDLC*GR
						GQQGLLVGWT**EQKRGQGKP
						QYSSHSSSNTLSSN/ASSSHSDD
					,	RWFDPLDPL/EPEQDPLS/KGCM
						SLAK/APRPAKPHKPPGSMGLC/
						GGGREAAGRSHHADRR/REVSP
						APAVAGQSKGYR/PKLYSSGSS
						TPTGLAGG/SRDPPRQPSTLWH
						RTWYL/YHTASAAVHRGLCRE
						LEQADQIPPSWYGRRPMGNS
4137	34505	В	4178	108	318	
4138	34506	Α	4179	103	540	RRGCESHKTLRRGTSWGLDAR
						GGGPGPGQVSAGRDGAEVWLS
						TCDRGHALSGSVEELLFLQN/G
						ARTER*EGPGEWPRPPPPGLASP
					}	ALWRFWAEQVGGSFQELESPS
						CRTARGSSRTWGSILQNSSWLF
						QDLGLHLAEGCFLETP
4139	34507	Α	4180	33	896	KITRHCTAPGKIRIVPKESQEST
						PQQDGAPGPGRATSCSARWSPR
		į				SWKSHELFCKMEPQVLEEPRA
						VLQDGAPGPGRATSCSARKGR
						GPEKPVQGLPN/GSVRAHSGGR
						AAPQPSPRGHGPGRG*TAAPLP
				ļ		HLCPLTPVLLQG*GPD*WPLGW
						ATMRPLPLRAQPAPPPPHWML
		İ			,	LTPSAPPPGTGKPQGGRGQTSG
						SCVPATDPHCRLSAPSPGKLGPP
						CDFLEPP*QRTTNWGSSEAGSP
						KSRCPRGHVPSGGSKGGIFLSL
						HFPGAPQSLEFPGSQPHGLIGAS
4140	34508	В	4181	1	625	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
4141	34509	Α	4182	160	1149	FASERMKVEPWRAGPGRRAWS
						EGAGQAPQKRARAGAEPQLPA
						TPALPGGKMVARRTKLA\RGTR
						RT\YPEPTV\YAAIPIKFSEKQQA
						SHYLYVRAHGVRQGTKSTWPQ
						KRTLFVLNVPPYCTEESLSRLLS
				}		TCGLVQSVELQEKPDLAESPKE
						SRSKFFHPKPVPGFQVAYVVFQ
						KPSGVSAALALKGPLLVSTESH
						PVKSGIHKWISDYADSVPDPEA
						LRVEVDTFMEAYDQKIAEEEA
						KAKEEEGVPDEEGWVKVTRRG
						RR\LCSPGLRQPACGCWRGRDG
		-				SAA\KRAAQLLRLAASREQDGA
						SSAA\RKKFEEDKQRIELLRAQR
						KFRPY
4142	34510	Α	4183	2	361	GTMVARRTKLA\RGTRRTGIPS
						PPC*A\AIPIMCSEKQQASHYLY
						GRAHGIQQGTKSTWPHKRTIFA
						FNGPPYCSEQESLSCLQSTCGL
						VQSVKLKEKLELGWESRSKFFH
						PKPVPVTEEQ
4143	34511	Α	4184	917	1128	
4144	34512	Α	4185	1	660	MAWQMMQLLLLALVTAAGSA
						QPRSARARTDLLNVCMNAKYH
						KTQPSPEDELYGQ/C/SWRKNA/
						CSFTSTTQEAHKN/TSHLYGFN
:						WNHCGEMVPACKRHFIQDTCL
						YE*PPNLGPWI\RRYAWLPGIQE
						LAEELNFPGVSAGSNPSSSSIQG
			:			WVPGILEPEPFFSTKISQVDQSW
						RKEWVLNVPLCKEDCEQWWE
						DCRTSYTCKSNGHKGWNWTSG
11.15	21512	 	1106	0.6	501	SNKCQVAAA
4145	34513	Α	4186	216	781	MDMAWQMMQLLLLALVTAA
			:			GSAQPR\SARA\RTDLLNVCMN
						AKHHKTQPSPEDEAVWPDP/W
]						MCKGSCRKTKSWNI/HRKSKCE
1						VGLA/WEACSVSAGTGRGPGC
				!		GRWVGAPQGP/CPRKCSSG*PT
						W/VQRSQNMEEMAVVNQSWR
						KERILNVPLCKEDCERWWEDC
						RTSYTCKSNWHKGWNWTSAPS
L					<u> </u>	AVCDPLL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4146	34514	IA	4187	3	625	QCRPWRKNACCSTNTSQEAHK
						DVSYLYRFNWNHCG\KMAPDC
						KRHFHPGTPALYE/CAPHNLGA
						WDPAGWIQSWRKERVLNVPLC
						KEDCEQWWEDCRTSYTCKSN
						WHKGWNWTSGFNK\CAVGAA
						CPTF\HFYFPTPTVLCNEIWTHS
						YKVSNYSRGSGRCIQMWFDAS
						PGATPIEEV\ARFY\VAAMSGAG
	}					PWAAWPFLLSLALMLLWLLS
4147	34515	A	4188	1	268	EQGRH/GSSTPVGPRGGPR\GAE
						HAPKHQGCGDRAGPQVGMDQ
						RRRDPPRAPAAPRPWCGQRAA
						LSSLGGSHLCDDA*VQPSAGLG
						KVLKF
4148	34516	A	4189	2	1632	WKRCPGLPRAAATFPSGSGAG
						GARREAGGRAPTPGPASPRTAR
<u> </u>						GHARNSPAPARTAGRTGSAGA
						WQTPCPAPLFPMSAGLPAACH
	1					WNPV*LRALKTG/LEGVLGGSA
						DTQHNRVTDGSLAPN\AACVYT
	1					PKINGNRHPNTCTKMFIVSLDA
						KGKKWKQPTVHQQRKRETCG
						LHPRKCLQYTPS*WSTTTGILPS
						RTPRISCVQFVKKKLGQAGLLG
						HPGACLLCTLP*\PAGVGTFLFP
						RGC*GVVH*LETHTCG
4149	34517	A	4190	2	87	
4150	34518	Α	4191	3	291	
4151	34519	Α	4192	112	286	AWLLWLTSLPWGSLYALALLA
						NKPAL*SLLLLRYTLLPPHHQC
	2.1522	 			500	EKVPRWNEPQPTLFP
4152	34520	A	4193	1	933	
4153	34521	В	4194	125	999	WWAL VECTI EL EUTEDEECE
4154	34522	Α	4195	135	1160	VWALVRSTLELFHTDDEEEGE
				i.		YDEVTEEVTEQVYLPAKAKVA
				ì		QEEEVHPYPSAPPHYYFEEKEW PDPPDLSFLEDTGRKVVAPVTE
						ł
		1	,			QHLERLLSVLFRQEFSRLDERD DAVEQLRGVCIRAWEKITSGGE
						QYPSFSAVKQGPKELYADFIAW
						` ` `
						NLLRQESLKKVISDSAAQDIVL
					1	QLLAFGNVNLDCQAALRPIRGK
						AHLVDYIKACDGIGAKQDSERF
						AFTIPVVNNLQPAKHFHYFTDG
						SSNGKASYSGSKGQNQQPIWIL
						SRHLKPYHEPDAKEEIPGG/CPR
						TPWLQPCRD*C*GGP*/PVTSNT
		1				R*TQPPTWGQIKKLSQMVEENL
<u> </u>			<u> </u>	<u> </u>	L	RKAGQLVTMTVYWN

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4155	34523	A	4196	502	578	LV*EDCIAERAEVLRNESYGIIID WSP*GMFSLNCTSQSACHGHT MFSW
4156	34524	Α	4197	2	408	
4157	34525	A	4198	3	853	LLKVIMSAKIFTKKNENSTERL CGDGEKRGPDFRTERSVWLLR LEEAVAMVQRGSRAPESRVVA QVLTLLDGASGDREVVVVGAT NRPDALDPALRRPGRFDREVVI GTPTLKQRKEILQVITSKMPISS HVDLGLLAEMTVGYVGADLTA LCREAAMHALLHSEKNQDNPV IDEIDFLEAFKNIQPS/LVFEASL GLMGIKPVDWEEIGGLEDVKPE VKTAH/WSLRQKSGHC/RSCAR LPTGLLATLGSGSGSGRATEAV
	1		_			SGPAG*KRASIGGSSQRPRRFPT
4158	34526	A	4199	266	370	AERINSITVFSETLKRFLQASGK *FHRDIHNSRN
4159	34527	A	4200	18	1780	MGDVNQSVASDFILVGLFSHSG SRQLLFSLVAVMFVIGLLGNTV LLFLIRVDSRLHTPMYFLLSQLS LFDIGCPMVTIPKMASDFLRGE GATSYGGGAAQIFFLTLMGVA EGVLLVLMSYDRYVAVCQPLQ YPVLMRRQVCLLMMGSSWVV GVLNASIQTSITLHFPYCASRIV DHFFCEVPALLKLSCADTCAYE MALSTSGVLILMLPLSLIATSYG HVLQAVLSMRSEEARHKAVTT CSSHITVVGLFYGAAVFMYMV PCAYHSPQQDNVVSLFYSLVTP TLNPLIYSLRNPEERSHRGVKL NECNQCFKVFSTKSNLTQHKRI HTGEKPYDCSQCGKSFSSRSYL TIHKRIHNGEKPYECNHCGKAF SDPSSLRLHLRIHTGEKPYECNQ CFHVFRTSCNLKSHKRIHTGEN HHECNQCGKAFSTRSSLTGHNS IHTGEKPYECHDCGKTFRKSSY LTQHVRTHTGEKPYECNECGK SFSSSFSLTVHKRIHTGEKPYEC SDCGKAFNNLSAVKKHLRTHT GEKPYECNHCGKSFTSNSYLSV HKRIHNRWI*/YYCRNFWRKAL IDLSSLR*FERAHTGYISYLLQH
		+		18		
4161	34529	A	4202	1	389	
4162	34530	С	4203	114	548	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4163	34531	A	4204	122	735	LRAQQQHN*VLTLHKPACTLST TS*K*LHKIRK*LWHLRDRAPFI FTSEMEYFITEGGK/NPQHFQDF VELCCRAYNIIRKHSQLLLNLLE MNSYNGYVGLLHNILQLEREG LATKEELQQNFPPLSVSLPFDQS INQISEHRSLIFNGQYPYGSCWF RQAVCKLIQKYAGEWGWIATA ELRAEIDLNVLKFTIQVLSWKV QASLQ
4164	34532	Α	4205	139	4496	KMAYSWQTDPNPNESHEKQYE HQEFLFVNQPHSSSQVSLGFDQI VDEISGKIPHYESEIDENTFFVPT APKWDSTGHSLNEAHQISLNEF TSKSRELSWHQVSKAPAIGFSPS VLPKPQNTNKECSWGSPIGKHH GADDSRFSILAPSFTSLDKINLE KELENENHNYHIGFESSIPPTNS SFSSDFMPKEENKRSGHVNIVE PSLMLLKGSLQPGMWESTWQK NIESIGCSIQLVEVPQSSNTSLAS FCNKVKK
4165	34533	A	4206		3150	MEKPRPLEAPSAWPQDDVQCG VTVGMDGAAVRANRTPWPQD LEQTKWIEIKKSAFTWSSQLSL NRGFLTCKDENNNAGLLRVSS YSSREDQLKNIASDSLFMLPGG LCQSPTGTSHCSNQMETQGQGS PGGAVRGDKALGPEKARQGCG MNGSGKYCKFRVLAIQGKPEC LATLMQPDLGDSPGLREMNVV EHLRASFPVEQWYWRGGQRGE AEGARSSKAENNTSLICNFRLD YAPIEKQWDLHFADYFAEDLK
4166	34534	Α	4207	1	1203	

KKKMYANNGAIDRKLLFEATF VTIEKCCDTNQKGDDTHALGQ PIRGHDKSLAGSFCYACRSEEG LSQYRAYDSRGQLIAVKDTQG HETRYEYNIAGDLTAVIAPDGS RNGTQYDAWGKAVRTTQGG\L TRSV\EYD\AGRVIRLTSENGS HTTFRYDVLDRLIQETGFDGRT QRYHHDLTGKLIRSEDEGLVTH WHYDEADRLTHRTVKGETAER WQYDERGWLTDISHISEGHRV AVHYRYDEKGRLTGERQTVHH PQTEALLWQHETRHAYNAQGL ANRCIPDSLPAVEWLTYGSGYL AGMKLGDTPLVEYTRDRLHRE TLRSFGRYELTTAYTPAGQLQS QHLNSLLTYRHANFAL 4168 34536 A 4209 757 907 RRYCRITVRWQSMWADNRIA VDAHYPYR*CRS\GRVTEKND\ LIPKG\VIRTUDERTHRYHYDSQ HRLVHYTRTQYAEPLVESRYL YDPLGRRVAKRVWRRERDLTG WMSLSRKPQVTWYGWDGDRL TTIQNDRTRIQTIYQPGSFTPLIR VETATAVMDRILKDHQIVVDIP HGEAWLRDDEERPMILLAGGTG FSYARSILLTALARNPNRDITIY WGGREEQHLYDLCELEALSIK HPGLQVVPVVEQPEAGWRGRT GTVLTAVLQDHGTLAEHDIYIA GRFEMAKIARDLFCSERNARED RLFGDAFAFI	SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
4167 34535 A 4208 I 1470 MLHSRGFLAEVFGILARHNISV DLITTSEVSVALTLDTTGSTSTG DTLLTQSLLMELSALCRVEVEE GLALVALIGNDLSKACGVGVEE GLALVALIGNDLSKACGVGVEE VFGVLEPFNIRMICYGASSHNI. CFLVPGEDAEQDGTGTSGIGAQ KKKMYANNGAIDRKLLFEATF VTIEKCCDTNQKGDDTHALGQ PIRGHDKSLAGSFCYACRSEEG LSQYRAYDSRGQLIAVKDTQG HETRYEVNIAGDLTAVIAPDGS RNGTQYDAWGKAVRTTQGGIL TRSVIEYDVAAGRVIRLTSENGS HTTFRYDVLDRIQETGFDGRT QRYHHDLTGKLIRSEDEGLVTH WHYDEADRLTHRTVKCGTAER WQYDERGWLTJUSHISEGHRV AVHYRYDEKGRLTGERQTVHH PQTEALLWQHETRHAYNAQGL ANRCIPDSLPAVEWLTYGSGYL AGMKLGDTPLVEYTRDRLHRE TLRSFGRYELTTAYTPAGQLQS QHLNSLLTYRHANFAL LIPKGVIRTDDERTHRYHYDSQ HRLVHYTRTGVAEPLVESRYL YDPLGRRVAKRVWRRERDLTG WMSLSRKPQVTWYGWDGRL TTIQNDRTRIQTIYQAGSTPLIR VETATAVMDRILKDHQIVVDIP HGEAWLRDDEERPMILIAGGTG FSYARSILLTALARNPNRDITIY WGGREEQHLYDLCELEALSLK HPGLQVVPVVEQPEAGWRGRT GTVLTAVLQDIGTLAEHDIYIA GRFEMAKIARDLFCSERNARED	NO:	1	hod	i	1		,
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DLITTSEVSVALICLDTTGSTSTG DTLLTQSLLMELSALCRVEVEE GLALVALIGNDLSKACGVGKE VFGVLEPFNIRMICYGASSHNL CFLVPGEDAEQDGTGTSGIGAQ KKKMYANNGAIDRKLFEATF VTIEKCEDTDTQKGDDTHALGQ PIRGHDKSLAGSFCYACRSEEG LSQYRAYDSRGQLIAVKDTQG HETRYEYNIAGDLTAVIAPDGS RNGTQYDAWGKAVRTTQGGIL TRSVIEYDIAAGRVIRLTSENGS HTTRRYDVLDRLIQETGFDGRT QRYHHDLTGKLIRSEDEGLVTH WHYDEADRLTTHRTVKGETAER WQVDERGWLTDISHISEGHRV AVHYRYDEKGRLTGERQTVHH PQTEALLWQHETRHAYNAQGL ANRCIPDSLPAVEWLTYGSGYL AGMKLGDTPLVEYTTRDRLHRE TLRSFGRYELTTAYTPAGQLQS QHLNSLLTYRHANFAL 4168 34536 A 4209 757 997 RRYCRITVRWQSM/WADNRIA VDAHYPYR*CRS/GRVTEKNDN LIPKGIVIRTDDERTHRYHYDSQ HRLVHYTRTQYAEPLVESRYL YDPLGRRVAKRVWRRERDLTG WMSLSRKPQVTWYGWDGDRL TTIQNDRTRIQTIYQPGSFTPLIR VETATAVMDRILKDHQIVVDIP HGEAWLRDDEERPMILIAGGTG FSYARSILLTALARNPNRDITIY WGGREEQHLYDLCELEALSLK HPGLQVVPVVEQPEAGWRGRT GTVLTAVLQDHGTLAEHDIYIA GRFEMAKIARDLFCSERNARED RLFGDAFAFI				İ	sequence		,
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WQYDERGWLTDISHISEGHRV AVHYRYDEKGRLTGERQTVHH PQTEALLWQHETRHAYNAQGL ANRCIPDSLPAVEWLTYGSGYL AGMKLGDTPLVEYTRDRLHRE TLRSFGRYELTTAYTPAGQLQS QHLNSLLTYRHANFAL 4168 34536 A 4209 757 907 RRYCRITVRWQSM/WADNRIA VDAHYPYR*CRS\GRVTEKND\ LIPKG\VIRTDERTHRYHYDSQ HRLVHYTRTQYAEPLVESRYL YDPLGRRVAKRVWRRERDLTG WMSLSRKPQVTWYGWDGDRL TTIQNDRTRIQTIYQPGSFTPLIR VETATAVMDRILKDHQIVVDIP HGEAWLRDDEERPMILIAGGTG FSYARSILLTALARNPNRDITIY WGGREEQHLYDLCELEALSLK HPGLQVVPVVEQPEAGWRGRT GTVLTAVLQDHGTLAEHDIYIA GRFEMAKIARDLFCSERNARED RLFGDAFAFI							QRYHHDLTGKLIRSEDEGLVTH
AVHYRYDEKGRLTGERQTVHH PQTEALL WQHETRHAYNAQGL ANRCIPDSLPAVEWLTYGSGYL AGMKLGDTPLVEYTRDRLHRE TLRSFGRYELTTAYTPAGQLQS QHLNSLLTYRHANFAL 4168 34536 A 4209 757 907 RRYCRITVRWQSM/WADNRIA VDAHYPYR*CRS\GRVTEKND\ LIPKG\VIRTDDERTHRYHYDSQ HRLVHYTRTQYAEPLVESRYL YDPLGRRVAKRVWRRERDLTG WMSLSRKPQVTWYGWDGDRL TTIQNDRTRIQTIYQPGSFTPLIR VETATAVMDRILKDHQIVVDIP HGEAWLRDDEERPMILIAGGTG FSYARSILLTALARNPNRDITIY WGGREEQHLYDLCELEALSLK HPGLQVVPVVEQPEAGWRGRT GTVLTAVLQDHGTLAEHDIYIA GRFEMAKIARDLFCSERNARED RLFGDAFAFI					[WHYDEADRLTHRTVKGETAER
PQTEALLWQHETRHAYNAQGL ANRCIPDSLPAVEWLTYGSGYL AGMKLGDTPLVEYTRDRLHRE TLRSFGRYELTTAYTPAGQLQS QHLNSLLTYRHANFAL 4168 34536 A 4209 757 907 RRYCRITVRWQSM/WADNRIA VDAHYPYR*CRS\GRVTEKND\ LIPKG\VIRTDDERTHRYHYDSQ HRLVHYTRTQYAEPLVESRYL YDPLGRRVAKRVWRRERDLTG WMSLSRKPQVTWYGWDGDRL TTIQNDRTRIQTIYQPGSFTPLIR VETATAVMDRILKDHQIVVDIP HGEAWLRDDEERPMILIAGGTG FSYARSILLTALARNPNRDITIY WGGREEQHLYDLCELEALSLK HPGLQVVPVVEQPEAGWRGRT GTVLTAVLQDHGTLAEHDIYIA GRFEMAKIARDLFCSERNARED RLFGDAFAFI							WQYDERGWLTDISHISEGHRV
ANRCIPDSLPAVEWLTYGSGYL AGMKLGDTPLVEYTRDRLHRE TLRSFGRYELTTAYTPAGQLQS QHLNSLLTYRHANFAL 4168 34536 A 4209 757 907 RRYCRITVRWQSM/WADNRIA VDAHYPYR*CRS\GRVTEKND\ LIPKG\VIRTDDERTHRYHYDSQ HRLVHYTRTQYAEPLVESRYL YDPLGRRVAKRVWRRERDLTG WMSLSRKPQVTWYGWDGDRL TTIQNDRTRIQTIYQPGSFTPLIR VETATAVMDRILKDHQIVVDIP HGEAWLRDDEERPMILIAGGTG FSYARSILLTALARNPNRDITIY WGGREEQHLYDLCELEALSLK HPGLQVVPVVEQPEAGWRGRT GTVLTAVLQDHGTLAEHDIYIA GRFEMAKIARDLFCSERNARED RLFGDAFAFI							AVHYRYDEKGRLTGERQTVHH
AGMKLGDTPLVEYTRDRLHRE TLRSFGRYELTTAYTPAGQLQS QHLNSLLTYRHANFAL 4168 34536 A 4209 757 907 RRYCRITVRWQSM/WADNRIA VDAHYPYR*CRS\GRVTEKND\ LIPKG\VIRTDDERTHRYHYDSQ HRLVHYTRTQYAEPLVESRYL YDPLGRRVAKRVWRRERDLTG WMSLSRKPQVTWYGWDGDRL TTIQNDRTRIQTIYQPGSFTPLIR VETATAVMDRILKDHQIVVDIP HGEAWLRDDEERPMILIAGGTG FSYARSILLTALARNPNRDITIY WGGREEQHLYDLCELEALSLK HPGLQVVPVVEQPEAGWRGRT GTVLTAVLQDHGTLAEHDIYIA GRFEMAKIARDLFCSERNARED RLFGDAFAFI							PQTEALLWQHETRHAYNAQGL
TLRSFGRYELTTAYTPAGQLQS QHLNSLLTYRHANFAL 4168 34536 A 4209 757 907 RRYCRITVRWQSM/WADNRIA VDAHYPYR*CRS\GRVTEKND\ LIPKG\VIRTDDERTHRYHYDSQ HRLVHYTRTQYAEPLVESRYL YDPLGRRVAKRVWRRERDLTG WMSLSRKPQVTWYGWDGDRL TTIQNDRTRIQTIYQPGSFTPLIR VETATAVMDRILKDHQIVVDIP HGEAWLRDDEERPMILIAGGTG FSYARSILLTALARNPNRDITIY WGGREEQHLYDLCELEALSLK HPGLQVVPVVEQPEAGWRGRT GTVLTAVLQDHGTLAEHDIYIA GRFEMAKIARDLFCSERNARED RLFGDAFAFI					:		ANRCIPDSLPAVEWLTYGSGYL
QHLNSLLTYRHANFAL 4168 34536 A 4209 757 907 RRYCRITVRWQSM/WADNRIA VDAHYPYR*CRS\GRVTEKND\ LIPKG\VIRTDDERTHRYHYDSQ HRLVHYTRTQYAEPLVESRYL YDPLGRRVAKRVWRRERDLTG WMSLSRKPQVTWYGWDGDRL TTIQNDRTRIQTIYQPGSFTPLIR VETATAVMDRILKDHQIVVDIP HGEAWLRDDEERPMILIAGGTG FSYARSILLTALARNPNRDITIY WGGREEQHLYDLCELEALSLK HPGLQVVPVVEQPEAGWRGRT GTVLTAVLQDHGTLAEHDIYIA GRFEMAKIARDLFCSERNARED RLFGDAFAFI							AGMKLGDTPLVEYTRDRLHRE
4168 34536 A 4209 757 907 RRYCRITVRWQSM/WADNRIA VDAHYPYR*CRS\GRVTEKND\ LIPKG\VIRTDDERTHRYHYDSQ HRLVHYTRTQYAEPLVESRYL YDPLGRRVAKRVWRRERDLTG WMSLSRKPQVTWYGWDGDRL TTIQNDRTRIQTIYQPGSFTPLIR VETATAVMDRILKDHQIVVDIP HGEAWLRDDEERPMILIAGGTG FSYARSILLTALARNPNRDITIY WGGREEQHLYDLCELEALSLK HPGLQVVPVVEQPEAGWRGRT GTVLTAVLQDHGTLAEHDIYIA GRFEMAKIARDLFCSERNARED RLFGDAFAFI							TLRSFGRYELTTAYTPAGQLQS
VDAHYPYR*CRS\GRVTEKND\ LIPKG\VIRTDDERTHRYHYDSQ HRLVHYTRTQYAEPLVESRYL YDPLGRRVAKRVWRRERDLTG WMSLSRKPQVTWYGWDGDRL TTIQNDRTRIQTIYQPGSFTPLIR VETATAVMDRILKDHQIVVDIP HGEAWLRDDEERPMILIAGGTG FSYARSILLTALARNPNRDITIY WGGREEQHLYDLCELEALSLK HPGLQVVPVVEQPEAGWRGRT GTVLTAVLQDHGTLAEHDIYIA GRFEMAKIARDLFCSERNARED RLFGDAFAFI							QHLNSLLTYRHANFAL
LIPKG\VIRTDDERTHRYHYDSQ HRLVHYTRTQYAEPLVESRYL YDPLGRRVAKRVWRRERDLTG WMSLSRKPQVTWYGWDGDRL TTIQNDRTRIQTIYQPGSFTPLIR VETATAVMDRILKDHQIVVDIP HGEAWLRDDEERPMILIAGGTG FSYARSILLTALARNPNRDITIY WGGREEQHLYDLCELEALSLK HPGLQVVPVVEQPEAGWRGRT GTVLTAVLQDHGTLAEHDIYIA GRFEMAKIARDLFCSERNARED RLFGDAFAFI	4168	34536	Α	4209	757	907	RRYCRITVRWQSM/WADNRIA
HRLVHYTRTQYAEPLVESRYL YDPLGRRVAKRVWRRERDLTG WMSLSRKPQVTWYGWDGDRL TTIQNDRTRIQTIYQPGSFTPLIR VETATAVMDRILKDHQIVVDIP HGEAWLRDDEERPMILIAGGTG FSYARSILLTALARNPNRDITIY WGGREEQHLYDLCELEALSLK HPGLQVVPVVEQPEAGWRGRT GTVLTAVLQDHGTLAEHDIYIA GRFEMAKIARDLFCSERNARED RLFGDAFAFI			1				VDAHYPYR*CRS\GRVTEKND\
YDPLGRRVAKRVWRRERDLTG WMSLSRKPQVTWYGWDGDRL TTIQNDRTRIQTIYQPGSFTPLIR VETATAVMDRILKDHQIVVDIP HGEAWLRDDEERPMILIAGGTG FSYARSILLTALARNPNRDITIY WGGREEQHLYDLCELEALSLK HPGLQVVPVVEQPEAGWRGRT GTVLTAVLQDHGTLAEHDIYIA GRFEMAKIARDLFCSERNARED RLFGDAFAFI							LIPKG\VIRTDDERTHRYHYDSQ
WMSLSRKPQVTWYGWDGDRL TTIQNDRTRIQTIYQPGSFTPLIR VETATAVMDRILKDHQIVVDIP HGEAWLRDDEERPMILIAGGTG FSYARSILLTALARNPNRDITIY WGGREEQHLYDLCELEALSLK HPGLQVVPVVEQPEAGWRGRT GTVLTAVLQDHGTLAEHDIYIA GRFEMAKIARDLFCSERNARED RLFGDAFAFI							HRLVHYTRTQYAEPLVESRYL
TTIQNDRTRIQTIYQPGSFTPLIR VETATAVMDRILKDHQIVVDIP HGEAWLRDDEERPMILIAGGTG FSYARSILLTALARNPNRDITIY WGGREEQHLYDLCELEALSLK HPGLQVVPVVEQPEAGWRGRT GTVLTAVLQDHGTLAEHDIYIA GRFEMAKIARDLFCSERNARED RLFGDAFAFI			l				YDPLGRRVAKRVWRRERDLTG
VETATAVMDRILKDHQIVVDIP HGEAWLRDDEERPMILIAGGTG FSYARSILLTALARNPNRDITIY WGGREEQHLYDLCELEALSLK HPGLQVVPVVEQPEAGWRGRT GTVLTAVLQDHGTLAEHDIYIA GRFEMAKIARDLFCSERNARED RLFGDAFAFI			ĺ				WMSLSRKPQVTWYGWDGDRL
HGEAWLRDDEERPMILIAGGTG FSYARSILLTALARNPNRDITIY WGGREEQHLYDLCELEALSLK HPGLQVVPVVEQPEAGWRGRT GTVLTAVLQDHGTLAEHDIYIA GRFEMAKIARDLFCSERNARED RLFGDAFAFI							TTIQNDRTRIQTIYQPGSFTPLIR
FSYARSILLTALARNPNRDITIY WGGREEQHLYDLCELEALSLK HPGLQVVPVVEQPEAGWRGRT GTVLTAVLQDHGTLAEHDIYIA GRFEMAKIARDLFCSERNARED RLFGDAFAFI							VETATAVMDRILKDHQIVVDIP
WGGREEQHLYDLCELEALSLK HPGLQVVPVVEQPEAGWRGRT GTVLTAVLQDHGTLAEHDIYIA GRFEMAKIARDLFCSERNARED RLFGDAFAFI							HGEAWLRDDEERPMILIAGGTG
HPGLQVVPVVEQPEAGWRGRT GTVLTAVLQDHGTLAEHDIYIA GRFEMAKIARDLFCSERNARED RLFGDAFAFI		İ	l				FSYARSILLTALARNPNRDITIY
HPGLQVVPVVEQPEAGWRGRT GTVLTAVLQDHGTLAEHDIYIA GRFEMAKIARDLFCSERNARED RLFGDAFAFI							WGGREEQHLYDLCELEALSLK
GRFEMAKIARDLFCSERNARED RLFGDAFAFI							
GRFEMAKIARDLFCSERNARED RLFGDAFAFI	1			1			GTVLTAVLQDHGTLAEHDIYIA
RLFGDAFAFI	-		1				GRFEMAKIARDLFCSERNARED
4169 34537 B 4210 1 3258	4169	34537	В	4210	1	3258	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
4170	34538	Α	4211	281	11571	CQCPGAACPTTSCRVIPHWA\Y
17170	34336	^	4211		1371	DEADRLT/HRTVN\GETAERWQ
						YDER/GWLTDISHISEGHRVA/V
						HYRYDEKGRLTGERQT/VHHPQ
						TEALLWQHETRH/AYNAQGLA
]	NRCIPDSL/PAVEWLTYGSGYL
						AGMK/LGDT/PAANLDIRIPYAT
						DPA/GNRLPDPELHPDSTLSM/W
ı						ADNRIARDAHYLYRY/DRHGRL
						TEKTDLIPEGV/IRTDDERTHRY
į						HYDSQ/HRLVHYTRTQYAEPLV
ĺ		l				E/SRYLYDPLGRRVAKRK/DRTR
						IQTMYQPGSFTPL/IRVETATGE
						QAKTQR/RQLADTLQQSDGED
				1		GGS\VVFP\PVL\VQML\DR\LESE
	}					SSADRVRSFISLANQSKCVEHA
	1	ĺ				Y*RWQCHLGVCWWSSHQIPAV
						PLTGSVTRWHCHLASSSEAGSV
	}			1		AWLPHLSEGHSNRTSSPELLRS
Ì						RMCAWHTLSAQSVHLVSLYIL
		ļ				EILALMNSINSL
4171	34539	Α	4212	311	788	
4172	34540	Α	4213	29	395	RIFHSVI\GVAAHKGGVYKTSVS
						VHL\AQDVAEIT/LLEGNDPQGT
İ						VS*YQGPGRTLIPLEAALRNIAH
	}					SLSIPPPKIFAAPTLRHYFALFFC
				1		GHSLFAPHIELLEAGTVLQLPQ
						GPWSSPTSF
4173	34541	A	4214	1	1033	MKMPEAIATKEKIDKWDLIKIK
						SFFSTPKETINRVNRHHTEWEDI
						SAIHLSDKGPISYIYKNLTRFTR
		ŀ				KKQPHYKVGKGNEQTRILESHP
						HLLKGLASTPFDSEGVRTERRD
1						IIKDGILTQWLLTSYSARKLGLK
	[STGHAGGIHNWRIAGQGLSFEQ
						MLKEMGTGLVVPGTAENARSC
						IRAYFYDIHETLCRQEEMALSV
						VDDHVREKLIWLRQHQEDMTI
İ						
		1				LLSEVSAACLHCEKTLQQDDCR
						VVLAKQEITRLLETLQKQQQF
						TEVADHIQLDASIPVTFTKDNR
						VHIG\PKM\EIRVVTLGIGMGAG
						KNLLSLF*V*NRVEFHGSPFPTI
1						WFLTWETVGFLK
4174	34542	В	4215	414	1022	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
4175	34543	Α	4216	896	1626	NATTIRYEHORVLKAAQYLHO
,						QGITRCNSSSTLTSAAPGRVDSP
ļ						PTSMIVAPALIIRFACFTASISEL
						CVPPSEKESGVTLRIPITHLMRE
						LTCRLIRDKSATVTGSHTLVVA
						RHCCYAAPGGCCLLANWLKPG
						LFGPIGVLSRRGTSVILPIGGFY
i						QWNPMICPNGVVPMQHG*RRG
						LAQERPLEEWLPVCRDMLNAF
						FLPDAETEAAMTLIEQQWQAII
						1
ŀ						AEGLGAQYGDAVPLSLLRDEL
İ						AQRLDQERISQRFLAGPVNICTL
ŀ						MPMRSIPFKVVCLLGMNDGVY
						PRQLAPLGFDLMSQKPKRGDRS
						RRDDDRYLFLEALISAQQKLYI
						SYIGRSIQDNSERFPSVLVQELI
						DYIGQSHYLPGDEALNCDESEA
						RVKAHLTCLHTRMPFDPQNYQ
						PGERQSYAREWLPAASQAGKA
		1				HSEFVQPLPFTLPETVPLKRYN
						DSVRAPTCAESRAIFTSTRNNTL
						QLFFNANFRRPWARGLATNVN
				1		DRRASVDHQIRMFHRIYQRVM
Ì		1				RATIRKGIRRDVEDPHYSSDAG
						TYLSPNSRQISNGNWQSHIGRSP
İ		l				SLLLCRTWGLLFTGKLVETRFI
ŀ		1				WPNRGVIPTGNERYIAHRRFLP
						MEPDDMPQWRCPHATWLAEA
						KMFDSLAKAGKYLGQAAKLMI
		l				GMPDYDNYVEHMRVNHPDQT
ŀ						PMTYEEFFRERQDARYGGKGG
4176	34544	Α	4217	838	1575	CFFLSPSPPSSPPSPNRSQTTEEE
,		l				TKRQE/ERERKREEEEEKKGRR
		l				KETKKRRNRQEGKQHRKEEKE
						GEKQTKQRTETERETKRRRENE
		l				QAKAHKGTRKRKEEQKKAKA
		ŀ				ARRRTHKRONPSRGREGTHPK
						QRQGKEE/VNRQNKEEAKQKR
						EEAGRTRR/EDRGRKDDKKERR
						QQQTEKKAKPKAEHGQERTDT
}						TTKKARQREGRPSERRREREE/
						MSKHDPQNRAEKTNEEKEEGR
						QHER*TQKSSTGI
L			l	l	<u> </u>	KILDK TOKSSTOI

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	1			sequence		
4177	34545	A	4218	Ti	692	MNALHAEALEMTSQFDQELAA
		'				KFEADHEMALLMNKDFDRDRE
	1					EQRRLAEQARREHEERIKREAA
						EQARRDAEAKHKAEIEAAARR
						EAEEKARAELAERQRIEAEQRA
						EREKKETEERARREKEEAVAAE
•	-			:		RRRQEEAEAARLVEEQRKAEE
	E					EARRAADKEHRRTVNRR/GLRR
						SDCSGHPRRIRTESSAGDRWRQ
						SAGRAHQILRQT*THTSLTTAS
						KNGAGLSSNSPTRKRS
4178	34546	Α	4219	3	1120	
4179	34547	Α	4220	1	831	MVKARKIMETPQPAWEMRVRI
[CTVDWSKLNPYIPDDFSLIKSEK
						KYDHPELIVDESNLRVVYAPSR
						YFASEPKADVSLILRNPKAMDS
						ARNQALLEGYFSFTATEDQLEQ
						AKSWYNQMMD/SPEKGKAFEH
						GNMPA\QMLLQVPYFLRE\ERE
						H*IIITPILHMRKQEQSG*/RNLP
						KAAQLSMMQDLQTLLMAASY
		ŀ				CSELGHVATQFQGMLACTRNP
						NSWDRNSETSGKAEGFIPMQLG
		ŀ				DVADPSVRCSSVSSLWGHSSPK
1100	24540		1001	ļ	1500	LLRSVCMANRICVKLQRWT
4180	34548 34549	A A	4221	1	1503 1113	
4182	34550	В	4223	1	760	
4183	34551	В	4224	1	1755	
4184	34552	С	4225	1	4215	
4185	34553	Α	4226	1	3240	
4186	34554	Α	4227	1989	2144	
4187	34555	A	4228	1	1203	
4188	34556	Α	4229	31	4767	EVENEDEDENIOUSEA EITH
4189	34557	Α	4230	31	512	EYRKSPDIRPIVIQHGEEAEITH
						HFR*QELADKTLIFEITHREMQR
						FQPVGTGDIREPVFVFFRWRLT NPFNILEHGEPEGIRVDAAVPR
						AVIGGLEDHIGVAVQKLQHKTF
						RYFPFIIQMVKDGVVPEGRPAF
4190	34558	A	4231	369	918	VHHLSLFLRIKILAHLTHNTQDF RPGMSNPWRDLFRTGVDPTND
130	134338		7431	309	710	RLSALVEI/YRMMRPGEPPTREA
1						AE/SLFENLFFSEDRYDLSA/VG
						RMKFNRSLLREEIE/GSGILSKD
1				-		DIIDVMKKL/IDIRNG/KGEVDDI
					1	DHLG/NRRIRSVGEMAENQFRV
						/GLVRCTGTVPPFLHQQKCEYH
						L/PQRPVASTLSRYF*HFRLRNF
						SMPRANEIKKGMV
	<u> </u>	<u> </u>	L	1	L	DIVIT KATALITAKUWI V

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence	ļ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
4191	34559	A	4232	3	1012	SVVLKIVERVDGYFPSPPLFLGA
						AFGSHVRSRALPDPCVLFRVTI
						YLCVLASVARTFSPLLPVSRKK
			}			HITPLGGFQLHETHLCLQDCRTI
1		İ				ILRPRLGEFGNKFSQLVDTIDLH
					}	\NQHGLAHL*KESARGQYSLLL
						LGMVSLCALILILWRVVYRSVT
		1				CPLADQTQALHRLLDGDIVSPF
		l				PETAGVRELDTIGRLMDAFRSN
						VHALNRHREQLAAQVKARTAE
1		l				LQELVIEHRQARAEAEKASQAK
						SAFLAAMSHEIRTPLYGILGTA
		1				QLLADNPALNAQRDDLRAITDS
						GESLLTILNDILDYSAIEAGG\K
		1				NVSVQSYVARLEPVAASGWHK
ŀ		1	•			YPWLN
4192	34560	Α	4233	1	502	
4193	34561	Α	4234	1	653	
4194	34562	Α	4235	2	300	YALATPPLSV/INQWQLALDKG
						QLPTF/VAGLAPQHPQYAAMHE
						SYWPYSALR/EILQRTGMLDGG
						PKITL/PGDDTPTDAVVSPSAVT
						NSHGR*VPTLGGVWGL
4195	34563	С	4236	40	105	
4196	34564	A	4237	355	526	
4197	34565	Α	4238	116	949	RPGTGRCSAVQLPVLLLRGPHS
						SHTVGTH\MVDLDSGQLCVYP
						GNSDESMPAATQARERLLADT
						AKKKAQIAELQSFVSRFSANA*
			ŀ			KSRQATSRARQIDKIKLEEVKA
						SSRQNPFIRFEQDKKLFRNALE
			ŀ			VEGLTKGFDNGPLFKTLNLVL\
						EVGENLPVLGTNGVGKSTL\LK
		ļ				TLVGDLHPDSGTVKWSENARI
		1				GYYAQDHEYEFENDLPVFEWM
		l				SQWKQEGDDEQAVRSILGRLLF
		l				SQGDIKKPAKVLSGGEKGRML
		<u> </u>				FGKLMMQKPNILIMDEPPTHP
4198	34566	Α	4239	1	319	MVKKMARAPMI\L\ALANPEPEI
						LACRHGRKEVRPDAIIC\TPGRS
						DYPNQSETNVL\CFPANVHRIPQ
						AASHLRAHQSRIPISLMSISAKIL
4199	34567	В	4240	263	1390	TYLLANQIQFLVKQH
4200	34568	A	4241	1	323	
4200	34569	A	4241	3	1855	
7201	127303	I.r	17272	1-	1000	<u> </u>

SEQ ID	SEQ ID NO:	L	SEQ ID NO:	4		Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/340,217	sequence	peptide sequence	detection, 1-possible nucleotide hiser tion)
		<u> </u>				
4202	34570	Α	4243	2	964	LLKHGVSLYIQDKEGLSALDLV
						MKDRPTHVVFKNTDPTDVYTW
						GDNTNFTLGHGSQNSKHHPEL
						VDLFSRSGIYIKQVVLCKFHSVF
	İ					LSQKGQVYTWGHGPG\GRLGT
]				,		WEMNRHAWVPRLVGRD*MVII
		1				VSPSWPAAKDHTVVLTEDGCV
						YTFGLNIFHQLGIIPPPSSCNVPR
		ł				QGLHNRQNRRYPPVPGSAGPTS
		ŀ				MEPSKIRPTGLKFSLTTQQQSEI
		l				DLGCSSLCWDYRREPLRLAYW
						FIKKDIAKDTDEETRRHGVSLYI
						QDKEGLSALDLVMKDRPTHVV
						FKNTGSLQFQSIPSCRESQILSEK
						QGDLFREEPMFGS
4203	34571	Α	4244	1	725	FRVDPRVRKHFGLFYAMGIVL
						MMEGVLSAC*HVCPNYSNFQF
ļ				1		DTSFMYMIAGLCMLKLYQTRH
						PDINASAYSAYASFAVVIMVTV
						LGVVFGKNDVWFWVIFSAIHV
						LASLALSTQIYYMGRFKIDLGIF
						RRAAMVFYTDCIQQCSRPLYM
		1				DRMVLLVVGNLVNWSFALFGL
						IYRPRDFASYMLGIFICNLLLYL
						AFYIIMKLRSS*KVLPVPLFCIV
						ATAGMWAC\ALYFFFQNLSSW
4204	34572	A	4245	1	833	MKPVWVATLLWMLLLVPRLG
						AARKGSPEEASFYYGTFPLGFS
		I				WGVGSSAYQTEGAWDQDGKG
						PSIWDVFTHSGKGKVLGNETA
						DVACDGYYKVQEDIILLRELHV
						NHYRFSLSWPRLLPTGIRAEQV
		1				NKKGIEFYSDLIDALLSSNITPIV
		1				TLHHWDLPQLLQVKYGGWQN
		1				VSMANYFRDYANLCFEAFGDR
		1				VKHWITFSDPRAMAEKGYETG
}						HHAPGLKLRGTGLYKAAHHII/
						KAHTL*VCFHAADKGIPETEKK
		<u> </u>				RRLNWTYSSTWLGRFHNHGRG
4205	34573	Α	4246	1	672	GTQNAVNG/VIIFLSWGDAVKS
						FWIYRGGRKREGPLFHA*Q\FLI
						YTIIIRAVGSIINYVIANYKLKFI
						TPGVIDFICTSLIAGILTIKLFLLI
						NQFEKQQIKKGRDITSARIMSRI
						IKITIIVGLVLLYGEHFGMSLSG
						LLTFGGIGGLAVGMAGKDILSN
						FFSGIMLYFDRPFSIGDWIRSPD
Ī						RNIEGTVAEIGWRITKITTFDNR
	<u></u>	<u> </u>				PLYVPNSLFSSISVENPG

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4206	34574	A	4247		347	PLRP\GQPGPGGAGT/RALRAPP LPSSSELCYGPGQPGRWPRCPQ FPSLPPHSS*LLHTGHWPCTLGY CFIPILRAAPPLPCKCASPVL\SC TYPLPAAPSLPVLVHTSIKCFCF HLQ
4207	34575	A	4248	43	446	VLPAVPRPGQ/PPCVPAPVQAPE PPRPSPGWSQATGSPGPAGAAP SWR\GLPAVPGHTAGVLGPGPP GQRQPGPGGAGTQLCGPHLFL RLLNVAYGPGQPAL/RNVPLQT APALTPSPQHLLCPFLSTHLLNA SVFIC
4208	34576	A	4249		1521	RIPESRLPTTAFVQAPWARSGSS GLRRWEKHAGQQVGWWARGP GVRGRQAAGGGAAALTCRGG AGSAVRSCAGLPSLASGSAGCR LHPSYSFGFKVGS/PTVPAALSS *STS/RGREHGGVTVVPVMTQN PRS\PDGPARVEDCEAIA*GTG WL\QQGIGTRPPGTGLGRAR\G APAVPQWNPV\KSCQGPGHPNR LPSHGPSSGEAGRGW/RGLQITP QL/PEVTHRRVLPGDHPATEA\G GFGTG*PGLPGRVPGPGVGGTY QAKALTPLGPVGLLAPASCAQC LQQSADGPGATGHL*ELAESQC RRQPTG\PPGQLAVSGWATVPG VPAAPRPFGPAQQPA/SVPTPSY WA/GSPGAAAWPESHRR*ACD WAW**VLPAVPRPGQ/PPCVPA PVQ\PQSHRGPAHDGARLY*GL PQAEQLHPGGGLPAVPGHTAG VLGPGPPGQRQPGPGGAGTQL CGPHLFLCFLNVAYGPGQPGR WPRCPQFPFLPPILSWNIFGVGT QKKKKKNQSFLKKKKKK
4209	34577	A	4250	167	582	RSLGLAVTEMVPWVRTMGQK LKQRLRLDVGREICRQYPLFCF LLLCLSAASLLLNR*RRSAEPGR RL/SL/LKVQTPGPCCLTVRRPRS CGTGDQWGERAPQRSGLGEHG GASRPEAQAGGVGLIASFPEAS SPELPFSHP

SEQ ID NO:	SEQ ID NO: of peptide	Met	SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence	1100	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4210	34578		4251	 402	1465	DI II CUDTA WETIVVVDVOLO
4210	34378	Α	4231	402	1403	DLILSHPTAWFTIKYKPKQLGL
	•					QELFPQGHSCAVCGKVKCKRH
						RPSLLLENYQPWLDLKISSKVE\ ESLSKDLELVLENFVYPWYRD
						VTDDESFVDELRITLRFFASVLI
						RRIHKVDIPSIYNQETIKSSNESI
		ŀ				*KWIVKARQKVKNTEFLQQAA
						LEEYGPELHVALRSRRDELHYL
						RKLTELLFPYILPPKATDCRSLT
					:	LLIREILSGSVFLPSLDFLADPDT
			į			VNHLLIIFIDDSPAEKATEPASTL
						VPFLQKFAEPRNKKPSVLKLEL
i i						KQIREQQDLLFRFMNFLKQEGA
						VHVL\QFCLTVEEFNDRILRPEL
						SNG*NAVSS/WKNCRRFIKHTV
			,			WMKVLTKLDLIPSLVEEIPR
4211	34579	Α	4252	1	1232	FPGRRFRLVVRLRGAEAASERQ
						VYSVTMKLLLLHPAFQSCLLLT
						LLGLWRTTPEAHASSPGAPAIS
						AASFL*DLIHRYGEGDSLTLQQ
						LKALLNHLDVGVGRGNVSQHV
						QGHRNPTTCFSSGDLFTAHNF\S
						EQLRIGSSELHEFCPTILQQLDS
						RACTSENQENEENEQTEEGRPS
						AVEVWGFGFLSVSLINLASLLG
				ļ		VLVLPCTEKAFFSRVLTYFIALS
						IGTLLSNALFQLIPERSYKNKAQ
						VDSLPTFLAQAGMLLWRVRIR
						RRVVDPIRESWMLPFTKIPLWG
						YGLLCVTVISLCSLLGASVVPF
						MKKTFYKRLLLYFIALAIGTLY
						SNALFQLIPENRRKWWQPVHN
	<u> </u>					TFGGSTAWHTDKSIEQSIDTLFD
						EVKKESEKETPSLQIGDLGPQES LKTFNNTNSPHH
4212	34580	Α	4253	3	924	VGACTAAARPLPIPQLQPILHHR
4212	34360	Λ	4233	٦	924	GEKSQLWAHSGSSWGFLAVAA
						VPPSHLCPPLQSRGWKRPP/PLA
						SAGVLPGCCCCACLVSPSLAQP
						AG\LGPKPAAPLGPGPWVSVAP
						CSRPGPCPGTRSPA/P*GHPAMG
						R\GVHEPRVGPAPPEKAIITETG
						AGLAERRGQGLGGGSSFRSAEP
						QGCRSLGPQSPGGDPAHTILRPP
						SQNGDCAEMHACRLHPAILGT
						HGTGGLAAQSHAPRALLPSCPS
						SQQPADGWCSLHLCLPGLLLAP
						RIHGPSTREGGPGHGTPGPTNP
						ASSGATRGTRRVRPSVPRSPTL
<u> </u>		Щ.	t <u> </u>	<u> </u>	<u> </u>	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
4213	34581	Ā	4254	1	318	VADPVGAARAGGQPLAGRVW
	:					PRAGGGHSPRVLGAAGPGPHV
ŀ	,					CTLRLPGTAIRTEPGAPLACAR
						AWPGSSPAG/PECLPTSC*P*EG
	•					QEPSSHVASLPWGPGVGETQ
4214	34582	A	4255	1	718	FFFFFSFLCHLYWVSPTPGPHG
						KLANMANWAPWPS*GLSKLVG
						KHSCPAG*LPGHARAQASGAP
						G/ISPDSSAREA*ECT/PCGPGPA
						APSTRGECPPPSSRPHS/SQQDP
						GRCSFAPAVPQDAGGQGHWCC
						APATGHSAPRGCPPARAAPTGS
			į			ATPAPPPAACAASSLSMVSAPS
						R*TTGIASSGTSIPETKHQGTPG
ŀ						TAPAGT/GPGGSTGPKA/PGPAP
						AHPTRLAGTSGHTAPPTCPPAV
4215	34583	Α	4256	702	1026	RSGRTQRAAGVSGSALHQVQS
						WP\HLKISADQRAGLLF*EHPFP
					!	PSASSGCLDVSISSYPVGSD*FIN
						GMARANGRWKTFTGLHSGKPL
						GFSDAFCQHHNLIILCWKTW
4216	34584	Α	4257	170	1049	RGSGCSAELVPSSRWRPGSRAG
						AAAGTETPG*PRVYVPAGNGE
						AGGPGAAWARRAAALPGTAA
						GPPRPAARPGAAPARGGPAPGA
						PAQALPR/TPTWPAAR*AQRAP
						SPPSWGS\AQPGHPGDLAAGVG
ŀ						RGAGGGHSRRGRHHHVRSLAD
						LLQLPGAAEGAGDRGHLPGPD/
						GERS*AASSFSAAGRAAGTASC
						CSAGGTPPSPCTILSTSSSSLAH
						VASSS/RRRAEGDTKVS/RGRAE
						GQDSETGREPGVLHRGSGRTQ
1						RAAGVSGS/RSAPSPVVATTSRS
		<u> </u>				LLTSVQGCFSENILSP
4217	34585	Α	4258	178	556	QSPQEHFHPECGRRDILCQVRQ
						EIRWPNPGEVHHLGLEICPVWI
						LQLHLALRTRAPEHPLQVHRPG
1						GGAV*RGVPPPLRLLQACDGPE
1						VPAAGRPRPARSSPGQWPP*/PA
				<u></u>		AVAPPVTERPPTPSAA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4218	34586	A	4259	5	1044	TGRILDGWHWAKELRLDCPLG
						DSRRPPFSRVSTEGSPAFLALRL
						PNVTAGS*EVSMLASTETPLVIT
						RPSPG\GHDPGPAPRGAAASPA
	i					GSPAP*QKSPRPLSAAAPPLLAS
						DPAPPRAAAPPADTESSVQPPA
:		İ				APHAGPWT\PSAPGPLDVHSPPP
				1		\PSRGPFVQSSAEPHNRPSPGAT
						RPRP\PPRGAAASPAGSPAP*QK
ļ						SPRPLSAAAPPLLASDPAPPRAA
						APPADTESSVQPPAAPHAGPWT
						LERSWAP*RSLPTPIPVADPLCR
						APLSHRYP*GDCQRSGLCHTSP
						GRASHLPGPGAHKRTPHACWL
		ŀ				PLECHRRSPHP*THPSG*PGPSP
						QSFFPEFLGSGP
4219	34587	Α	4260	2	576	CLVNSTTRRSFQLRLVPVPKFQ
						PPHMTVR*LFNFGRQLTATTFS/
						LRKSYAVREAYELQNCPDPPPF
		l				QNGYMINSDYSVGQSVSFECYP
						GYILIGHPVLTCQHGINRNWNY
						PFPRCDAPCGYNVTSQNGTIYS
						PGFPDEYPILKDCIWLITVPPGH
					:	GVYINFTLLHTEAVNDHIAVW
						YENLSSQNICDCDQQF
4220	34588	Α	4261	1	837	MWAGNAWRAALSGVPCGRSA
						QSVLAQLRGILEGELEGIRGAG
		l				TWKSERVITSRQGPHIHVDGVS
						GGILNLTSVRFIRGTQSIHKNLE
		1				AKIARFHQREDAILYPSCCDAN
		l				AGLFEVLLRPEDAVLSDELNCA
		l				SIIHGICLCKAHKYHYCHLDVA
						YLETKLQEAQKHRLFLVATDG
				}		AFSMDGDIVPLQKICRLASRYG
		1		1		ALVFVDECHATGFLGLTGQGT
						DELLGVMGQVTIINSTLGKALG
						GASGGYTTGPGPLVSLL/RAQP
				i		YLFSNSLPPAVVGCTSKAL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		ļ		sequence		
4221	34589	Α	4262	1	2142	MIILIDAEKAFDKIQQPFMLKTL
		ŀ	-			NKLGIDGTYLKITRAIYDKPTA
		!				NIILNGQKLEAFPLKTGTRQGCP
						LSPLLFNIVLEVLAQAIRQEKEI
						KGIQLGKEEVKLSLFADDMILY
						LENPIVSAQKLLKLISNVSKVSG
						YKINVQKSQAFLYTNNRQTESQ
						IMSEFPFTIATKRIKYLGIQLTRD
						VKDLFKKYKPLLNKIKEDTNK
						WKNIACSWIGRINIMKMAFPR
						WELNNENTWTQEGEHHTLGPV
		ŀ				VGWGKRGGIALVDIPNVNDKL
						MVLEVLARAIRQKKEIKGIQLG
						KEEVKLSLFADDMIVYLENSIV
						SAQNLKLISNFSKVSGYKINVQ
						KSQAFLYTNNRQTESQIMSEFP
						FTIATKRIKYLGIQLTRDVKDLF
						KENYKPLLKEIREDTNKWKNIP
	1					CSRIGRINIMKMAILPKVIYRFN
						DIPIKLPMTFFTELEKTTLKFIW
						NQKRACIAKTILSKKNIAGGITL
]					PDFKLYYKATVTKTAWYWYQ
	1	ŀ				NRDIDQWNRTEASEVTSHIYNH
	1	ŀ				LIFYKPDKNKKWGNDSLFNKW
	1	ļ				CWENWLAICRKLKLDPFLTPYT
						KIHSRWIKDLNVRPKTIKTLEEN
	ļ					LGNTIQDIGMGKDFMTKTPKA
						MATKAKVDKWDVIKLKSFCTA
						KETTIRVSRQPTEWEKIFAIYPS
	ļ	ŀ				DKGLISRIYKELKQIYRKK\TNN
	ļ					PIKKWAKNMNRHFSKEDIYAA
	<u>.</u>					NRQMKKCSSSLVIREMQIKTTM
4222	34590	Α	4263	1	1989	
4223	34591	Α	4264	1	1104	
4224	34592	Α	4265	1	879	
4225	34593	Α	4266	1	1659	
4226	34594	В	4267	1	1500	
4227	34595	В	4268	1	1962	
4228	34596	В	4269	1	1716	
4229	34597	Α	4270	1	1152	
4230	34598	Α	4271	1	4752	
4231	34599	Α	4272	1	2790	
4232	34600	Α	4273	1	3477	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
4233	34601	A	4274	1	1007	MLDASCHRTSDSKFFSFGVQTG
4233	134001	<u> </u> ^	42/4	I	1007	FLTPELAHLVGPCDRDHNSSPA
						REQNWTENEFDELTEVGFRKW
		İ				VITNSSELKEHVLTQSKEAKNL
						EKRAIKQEKEIKGIQLGKEEVK
	:					LSLFADDMIVYLENPIVSAQNL
						LKLISKFSKVSGYKINVQKSQA
						LLYTNNR\SQIMSELPFTIAMKR
						IKYLGIQLTRDVKDLFKDNYKP
			1			LLKEIREDTNKWKNIPCSWLGR
						INIMKMAILPKAIYRFNAIPIKLL
						*TFFTELEKTTLNFIWNQKRARI
	,					AKTILSKKNKAGGITLPDFKLY
1						HKATVTKTAWYWYQNRYIDQ
			:			WNRTEASEITPHIYNHLIFDKPE
4234	34602	A	4275	737	2460	RIKYLRIQLTRDVKDLFKENYK
4234	34002	A	42/3	137	2400	SLLNEIKEDTNKWKNIPCSWIG
			1 1			RMNIIKMAI/LPKVIYRFNVIPIK
						LPMTFFSELEKSTLKFIWNQKR
						ARIAKTILSQKNKAGGIMLPDF
						KLYYKATVTKTAWYWYQNRD
						IDQWNRTEPSEMTPHIYNHLIFD
						KPDKNKQWGKDSLFNKWCWE
			1			NWLAIGRQLKLDPFLTPYTKIN
						SRWIKDLNVRPKTIKTLEENLG
						NTIQDISMGKDFMSKTPKAMA
						TKAKMDKWDLIKLKSFCTAKE
						TTIRVNRQPTEWEKNFAIYSSD
			:			KGLISRIYKQLKQIYKKKTNNPI
						KKWAKDMNRHFSKEDVYAAN
						RHMKKCSSSLAIREMQIKTIMIY
1		Ì				HLTPVTMAIIKKSGNNRCWRG
						CGEMGTLLYCWWDCKLVQPL
						WKTLWQFLRDLELGIPFDPAIP
						LLGIYPKDYKSCCYKDTCTPKL
						ARDDQIHILKQHRRKELETRQK
						QYRAWYEINPFHSVWPVTAGK
						SPRHQLPVWVHNPQTSPYLQL
						QTRDGEESNENNFGSTILASDFF
						AEIDKLSILQIHMEMEGTQNSQ
						NNLDKKKTKMEDLHFSISKLLH
						SYSIQDNVISA
4235	34603	Α	4276	3	355	RQPVHLVHELPQQSWGICLNSS
						EQHGALQHSSLHL/RMCSEPWS
						SADPQ*R*TCRNL*LPVRGPPRR
	:			ł		TDLFSVSSKSTLKEWPLLLMIL
						AELGSYLILSGRREESYFTSLVL
						ISIGDC
4236	34604	В	4277	78	791	
	I	Ь	I		L	<u> </u>

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	location of first codon for peptide	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
				sequence		
4237	34605	A	4278	1	3395	MIISIDTENAFEKIQQPFMLKTL
ŀ						NKLGIDGTYLKIIRAIYDKPTAN
		l				IILNGQKLEAFPLKTGTRQGCPL
						SPLLFNIVLEVMARVIRQEKEIK
		ŀ				GIQLGKEEVKLSLFVDDMIVYL
						ENPIVSAQNLLKLISNLSKVSGY
						KINVQKSQAFSYTNNRQTESQI
						MNGLPFTTASKRIKYLGIQLTR
						DVKELFKENYKPLLNEKKVDT
						NKWKNIPCSWIGRIN\ILKMAIL
						P/KELEKTTLKFIWNQKRACIAK
						SILSKKNKAGGITLPDFKLYYK
						ATVTKTAWYWYQNRDIDQWN
						RTEPSEII\PHIYNHLIFDKPDKN
						KKWGMGSLFNKWCWENWLAI
						CRKLKLDPFLTPYTKINSRWIK
1	,					DLNVRPKTIKTLEENLGNTIQDI
		ŀ				DMGKDFMSKTPKAMATKAKID
		ŀ				KWDLTKLRSFCTAKETTIRVNR
						QPKEWEKIFAIYSSDKGLISRIY
		ľ				KELKRIYK/KKNNPIKKWAKD
						MNRYF*KEDIYAANRHMKKCS
						SSLAIREMQIKTTMR/YHLTPVR
						MAIIKKSGNNRWEMNNENTWT
						QEGEHHTLG/HC/WWKARRSRS
						CLTWMAAGKKRMRKTLQMT
4238	34606	В	4279	1	2011	

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SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			:	sequence		
4239	34607	A	4280	1	2661	MTMNFVADSHTGRNPLASAAG
						AKTGLRPLPRPCGARVWNPPD
						AGGGGVGSLKTSTPLGPLSAAN
						SPVHQGSVPQTRARGGGTLFQE
						VVTSRTLAFRNSLSAFTEVTSG
						TVSGRKGGRSTHLAGRRVSGG
						EGSRKAAAAALAAVAAAPGPV
						RRCSSQSCFSSSGSSHYSARTSP
İ	İ	ļ				VRVRPRRSLSSRSAAGNRAEAT
ļ		l				ESAMEKTLETVPLERKKREKEQ
				,		FRKLFIGGLSFETTEESLRNYYE
						QWGKLTDCVVMRDPASKRSRG
						FGFVTFSSMAEVDAAMAARPH
						SIDGRVVEPKRAVAREESGKPG
						AHVTVKKLFVGGIKEDTEEHHL
						RDYFEEYGKIDTIEIITDRQSGK
						KRGFGFVTFDDHDPVDKIVLQK
						YHTINGHNAEVRKALSRQEMQ
						EVQSSRSGRGDGYGSGRGFGD
						GYNGYGGGPGGGNFGGSPGYG
		ŀ				GGRGGYGGGGGPGYGNQGGGY
		ŀ				GGGYDNYGGGNYGSGNYNDF
		ŀ				GNYNQQPSNYGPMKSGNFGGS
						RNMGGPYGGGIWKNTSITERK
į		İ				KSRKLDLIQSKKGSRTKEAPQP
Ę		ŀ				PVASLCMHLGHWSRLMVSPGA
		ļ				QLTGKNSHGLSVSSVRKSNVGP
						RRLCAAMKATGPDNAQSQVSP
						PGHAPSAEDPTGSRTVSSPCTD
						RPHPFLSRPKPPTQISLVLPLKT
					1 .	DGALERMPQQL/HIASS/GAKVP
						NPSTQTPPVLLAFFYPFNLPP*N
4240	34608	Α	4281	1	908	MRKVKGKNRQSFKCLPPPSGA
1						LQAHGAAAPHGSLLTLHLHLV
						PVSSAAMKATGPDNAQSQVSP
						PGHAPSAEDPTGSRTVSSPCED
						RPHPFLSWPTWISLALLLKTDG
		l				ALERMPQQLPSLHPSQGTQSIH
						PDPSSTSSFLLPFQPPTLKRAAFP
						CPPSIVNPAVWDTSTPSVAEHH
						TPIRITLKEPTQFLSQKQYPIPQA
]				ALVGLQPIISHLLASHLLRPTDS
						PFNTPILPVKKPNGTYRLVQDL
						RLINQAVLPLVQE/DYSVLLYLP
						LNVTPGLPPATAFSYPPSPGPVA
						RARLASRLHSHAA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
Ì	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
4241	34609	Α	4282	1	915	MPNYVTFTDTKQLISDTPNNQV
						PMNRASMAFDAKCLTGCRFDD
						AIVQFDMTYWPFTVVNDAGRP
				}		KVQVEYERDKKLLPIGGVFYGS
1						DKDEGNCKSLPWEDCYQCCET
				ŀ		SQNVQDFLFLDVTPLSLDIKTA
ĺ		1				DGVMAVLIKCDATIPTRQTQTF
Ī	ļ					TTYSDKPSM/LIAKDKNLLRKFE
		1				LTGVPPAHHGAHQIEVTFDINA
		l				KGILNV\TLTDDKGHLSKEDIEP
		l				MVQETEKYKAEDEKQRDKVSS
		İ				KNSLDPYVFNMKATAEDEKLQ
						VKINNEHKQKILSKCHEIINWL
		ŀ				DKNQTAEKEEFEHAQQELEKSS
4242	34610	Α	4283	1	994	MHQTKKGNQWHFGMKAHIGV
		1				DAKSGLTHSLVTTRPNEHDLNQ
Ì	:					LGNLLHGEEQFVSADAGYQGA
		1				PQREELAEVDVDWLIAERPGK
						VRTLKQHPRKNKTAINIEYMKA
						SIRARVEHPFRIIKRQFGFVKAR
						YKGLLKNDNQLAMLFTLANLF
1		1				RADQMIHCTRGEGLITTKIPKAP
i		1				DNGSYCLPSKNDDSEEEDPEMS
		ł				PMVVTKMKEIAEAYLGKTVTN
l		ł				AVLTVPAYFNDSQRQAT/KKDA
						RTIAGLNGLRISNEPTAAAIAYG
						LNQKVGTERNVLIFDLGGSITPR
	1					IRTPETGSDDAIKSILEQAKKEIE
						SQKGGECDPCRQSLRPPGPAAN
4243	34611	Α	4284	3	677	
4244	34612	Α	4285	30	365	EEAETVLVGQLKQLSSCLAVH
		ļ.				KYRPETKQEKKQRLLARAEKK
						AAGKGDVPTKRPPVLRAGVNT
						VTTLVENKKAQLV\CRKMGVP
						YCIIKGKARLGRLVHRKTCTTV
						AFTQVN
4245	34613	Α	4286	3	432	NSRVDDFVAAQDAKGKKVAP
		Ī				APAVVKKQEAKKVVNPLFEKR
						PKNFGIGQ\QRLLARAEKKAAG
						KGDVPTKRPPVLRAGVNTVTT
						LVENKKAQLVVIAHDVDPIELV
1						VFLPALCRKMGVPYCIIKGKAR
	<u> </u>					LGRLVHRKTCTTVAFT
4246	34614	С	4287	62	217	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4247	34615	Α	4288	2	801	PKGKKAKGKKVAPAPAVVKK
						QEAKKVVNPLFEKRPKNFGIGQ
						DIQPKRDLTRFVKWPRYIRLQR
						QRAILYKRLKVPPAINQFTQAL
						DRQTATQLLKLAHKYRPETKQ
						EKKQRLLARAEKKAAGKGDVP
						TKRPPVLRAGVNTVTTLVENK
						KAQLVVIAHDVDPIELVVFLPA
						L\CRKMGVP\YCIIKGKARLGRL
						VHRKTCTT\VAFTQVNSEDKG\
						ALAKLVEAIRTNYNDRYDEIRR
						HWGGNDLRPK\SVARIAKLEKA
	<u> </u>					KAKELATKLG
4248	34616	В	4289	1	273	
4249	34617	Α	4290	1	441	
4250	34618	В	4291	47	482	
4251	34619	Α	4292	1	762	
4252	34620	Α	4293	1	890	MSKSESPKEPEQLRKLFIGGLSF
		ļ				ETTDESLRSHFEQWGTLTDCVV
		İ				MRDPNTKRSRGFGFVTYATVE
		ŀ				EVDAAMNARPHKVDGRVVEP
						KRAVSREDSQRPDYFEQYGKIE
						VIEIMTDRGSGKKRGFAFVTFD
		ŀ				DHDSVDKTVIQKYHTVNGHNC
						EVRKALSKQEMASASSSQRGRS
		ŀ				GSGNFGGGRGGGFGGNDNFGR
		İ	`			GGNFSGRGGFGGSHGGGGYGG
						SGDGYNGFGNDGSNFGGGGSY
		İ				NDFGNYNNQSSNFGPMKGGNF
		ļ				GGRSSGPYGGGGQYFAKPRNQ/
		<u> </u>				GGYGGSSSSSSYGSGRRF
4253	34621	Α	4294	1	1674	
4254	34622	A	4295	1	506	KYHTVNGHNCEVRKALSKQEI
						ASASCSQRGRSGSGNFGGDRG
						GGFGGNDNFGRGGNFSGHGGF
						GGSCCGGGYGGSGDGYNGFGN
						DASNFGGGGS/YNEFG/NYNNQ
						SSHFGPLS/GGNFGGRSS/SPLGG
						APASTYVKGPNSQRTQNEGWF
1						EG*APWRGDGGARGNKGGGA

SEQ ID NO:	SEQ ID NO: of peptide sequence	ı	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4255	34623	A	4296		1445	MKCLKFINHKEILEASERKQAE SLDFPFKKLRWHLCEGWIEEER DESRKSETIFKDLFKVPVLKETI YYKFYGPPVYQIETVYFMALSP PKSKQFDKTKQNNNNKKTHQF VIVFFKTDEHLSARGRRRRSIVK VSLLPAVIGLKSKFLKKPDQLR KLFI\GG\LSFETT\DESLEEPFSR QWGKRYTDSVVMRDPNTKRSR G\FGFVTYAT\VEEVDAA\MNA RPHKVEWKELLEPKRA\VSRED SQRPGCPH*LVKKIFVGGIKEDT \EEHHLRDYFEQYGKIEVIEIMT D\RGSGKKRGFAF\VTFD\DHD\S VDKIVI\QKYHTVNG\HN\CEV* KSPVKSKKMASASSKPKEGRSG FWETFGGGSWEVGFGGN\DNF GRG\GNFSWSVVAFGGSRG\GG GYG\GSGDG\YNGFGNDG\SNF
4256	34624	A	4297	1	920	GPMKGG\NFG\GRSSGPYG\GGG QYF\AKPR\NQGGYGGSSSS\SN DPGDTPNTASAPNCRSGKGRSS
						SPEHIPPLEKLEDSMQTNPSTNP EPGRLAEWLDPEERQQSLQFGL QEATSIGKGGQYYIKGTPHGTK ESEQQPSALDLPSDRAYPNEKE PENQLWRLVIKLIKEAPEKGAY LNVIKAVYDKPTNGEKLRAFPL RTGTKHRCPLSPLLFNILLEVLA RAIRQEKEIKSIQIGKEEVKLSLF ADDIIIYLESPKYSSRKLQELIKE FSKVSRYEINVHKSVALLYT\NS NQAENQIKNSASFTIAAKNKIK YLGIYLTKDAKDGYKENYKTL MKEIIDDKNKQKYIP
4257	34625	Α	4298	1	1194	
4258	34626	Α	4299	3	1834	
4259	34627	A	4300	285	502	
4260	34628	В	4301	77	1306	
4261	34629	A	4302	1	354	
4262	34630	Α	4303	1	1182	
4263	34631	В	4304	1	1995	
4264	34632	В	4305	1	1518	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	T F	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4265	34633	A	4306		918	MCPVGPWTHPVVISPVSECIVGI DILGSWQNLHIGSLTDITMVHYI DDMMLIGSSEQEVANSLDLLV RHLHARGCKINPTKIQGTSTSV KFLEFQWCGVCQDIPSKWVLE QKALQQVQAAVQAALPLEPYD PADPMVLEVSVADGVAVWSL WQAPIARIHGSRNQGVEVEVSP LTNIPSDPLAKFLFPAPSTLCSA GLELLVPEGGTLLPGNTTMIPL NWKLRLVPGYFGLLLALSPQA KNGVTVLAEVIDPDYQDEITLL FHNGGGEEYARNTGDPLRHLL VLPSPMIKVNGK\LQHPNPGRT
4266	34634	В	4307	1	1599	
4267	34635	В	4308	1	1569	
4268	34636	Α	4309	3	422	
4269	34637	Α	4310	1	1089	
4270	34638	A	4311	2	549	LKMTAMQRPMEKRMMNREIIL KERLSLTGIDIKILKKRSIMKVE SHRGEQISVSSLALQRIKYLGIQI TRDVKDLFKENYKPLLNKLKE DTNKWRNVPCPRVGRISIVKM AILPK/ILKKKTTLKFIWNQKRA HIAKTILSKKNKAGGITLPDFKL YYKAT/KTAWCWYQNRDTDQ WNRTKPSEI

472

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4271	34639	A	4312	371	3036	LIAYQPKKVQDQMDSQPNSTR VLEVLARAIRQEKEIKGIQLGK/ EEVKLSMFADDITAYLENPIVS APNLLKLISNFSK/VSGYKINVQ KSQAFLYTNKRQTE\QIMSELPF TVASKRIKYLGIKLRRDVKDLF KENYKPLLNEIKEDTNKWKNIP CSWIGRINIVKMTILPKVIYRFN AIPIKLPMTFFAELEKTTLKFIW NQKRAHIAKTILSQKNKAGGIM LPDFKLYYKATVTKTAWYWY QNRDIDQWNRIEPSEIIPHICKH LIFDKPDKNKKWGKDSLFNKW CWENWLAICRKLHLDPFLTPYT
						KINSRWIKDLNVRHKTIKTLEE NLGNTIQDIGMGKDFMTKTPK AMATKAKIDKWDLIKLKSFCT AKETTIRVNRQPTEWEKIFATY SSDKGLISRIYNELKQIYKKKTN NPIKKWAKDMNRHFSKEDIYA AKKHMKKCSSSLAIRETLYNDR RIGKLTQTCDETAFQPHVCTISR PMLSSPYRSSLTEKWSQDFSKP PYPFLFHKGYLNPREQDKEVLT RAIRQEKERKGIQLGKEEVKLS LFADDMIVYLENPIVSAQNPLK VVSNFSKVSGYKISVQKSQAFL YTNNRQTESQIMSELPFTIASKR IKYLRIQLTRDVKDLFKENCKP LLNEIEEDTNKWKNIPCSWIGRI NIVKMAILPKVIYRFNAIPIKLP MTFFTVLEKTTLKFIWNQKRAH
4272	34640	В	4313	1	1995	
4273	34641	Α	4314	3	549	
4274	34642	A	4315	3	614	EAYGQTECTGGCTFTLPGDWT S\GQFINILEMCLELSPCKSFSAD SARYVLGHVGVPLACNYVKLE DVADMNYFTVNNEGEVCIKGT NVFKGYLKDPEKTQEAL\DSDG WLHTGDIGRWLPDIENHNRLIV CTLTNTSWRSHKIIVLKTYQKA DDTKTPKETTFQNIMNLFLKEK RATAVLIRGGVGETSTDLSKKK PAKLLANF
4275	34643	A	4316	1	478	MKLDLHLSPYTKINSRWIKDLN LRPETIKILEDIIRKTLLDIGLGK DFMIKNPKVNATKTKINKWDLI KLK\NFCTAKEISSREIREPTEW EKIFANSASDKGLISRIYKELKQ IRSTLQLLFGISELPASLFLGFGA IMKSKKASLNTSTAILRQLIW

NO: of peptide sequence nod	SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
125	NO:		ı	1		codon for last amino acid	
4276 34644 A 4317 I 1125 MCHGIGAQIIPSHQTVQLDITAF LETVKKNKHKFYPAFIHILARL MNAHPEFRAMMKDGFIENMFF VSANPWSFSFDLNVANMDH FFAPVFTMGKYYTQGBKVLMP LAIGGPLESPDRDGGPLESTNR DASPESWSCRKSTFRLVAWYS AAKVFIRDKLMERNRRTGTR EKARIWEVTDRTVRTWIGEAV AAAAAAGOGGFRVDLARRSIR DENARSQNPVHTEGDMNMNIK KIVKQATVLTFTTAFLAGGATQ AFAKENNQKAYKETYGVSHIT RIDMLQIPKQQQNEKYQVPQF DQSTIKNIESAKGLDVWDSWPL QNADGTVAEVNGYHVVFALA GSPKDADDTSIYMFYQKVGDN SID\SWKNAGRVF 4277 34645 B 4318 I 1374 4278 34646 A 4319 I 1293 4279 34647 A 4320 I 1278 4280 34648 A 4321 I 1254 MNNMIKKIVKQATVLTFTTALL AGGATQAFAKENNQKAYKETY GVSHITRHDMLQIPKQQNEKY QVPQFDQSTIKNIESAKGLDW DSWPLQNADGTVAEYNGYHV VFALAGSPKDADDTSIYMFYQ KVGDNSIDSWKNAGRVFNOW DSWPLQNADGTVAEYNGYHV VFALAGSPKDADDTSIYMFYQ KVGDNSIDSWKNAGRVFNOW DSWPLQNADGTVAEYNGYHV VFALAGSPKDADDTSIYMFYQ KVGDNSIDSWKNAGRVFNOW DSWPLQNADGTVAEYNGYHV VFALAGSPKDADDTSIYMFYQ KVGDNSIDSWKNAGRVFNOSD KFDANDPILKDOTGEWSGSATF TSDGKIRLFYTDYSGKHYGKOS LTTAQVNVSKSDDTLKINGVNV GYYCEFSLETNKALYYGGGTNFF RKESQKLQQSAKKRDAELANG ALGIIELNNDYTLKKVMKPUTL TTDDFERANVFKMNGKWYL FTDSRGSKMTIDGINSNDIYML GYYSNSLTGPYKPLNKTGLUVL MGLDPNDVTFTVSHFAVPOAK GNNVVITSYMTNRGFFEDKKA TFAPSFLMNIKGNKTSVVKNSIL EQGQLTVN		sequence		09/540,217	1 ' '	of peptide sequence	deletion, \=possible nucleotide insertion)
LKTVKKNKHKFYPAFIHILARL MMAHPERMAMKDGFIENMF		:			sequence		
MNAHPEFRMAMK DGFIENMFF	4276	34644	A	4317	1	1125	MCHGIGAQIIPSHQTVQLDITAF
VSANPWVSFTSFDLNVANMDN FFAPVFTIMGKYYTQGDKVLMP LAIGGPLESPDRGGPLESTNR DASPESWSCKKSTPRLVAWVS AAKVFIRDKLMERRRRGTGRT EKARIWEVTDRTVATIGEAV AAAAADGGGFRVDLARRSIRK DRNARSQNPVHTEGDMNMNIK KIVKQATVLTFTTAFLAGGATQ AFAKENNQKAYETYGVSHIT RHDMLQIPKQQONEKYQVPQF DQSTIKNIESAKGLDWDSWPLQNADGTVAEYNGYHVVFALA GSPKDADDTSIYMFYQKVGDN SID\SWKNAGRVF		ŀ					LKTVKKNKHKFYPAFIHILARL
FFAPVFTMGKYYTQGDKVLMP							MNAHPEFRMAMKDGFIENMFF
LAIGGPLESPDROGGPLESTNR							VSANPWVSFTSFDLNVANMDN
DASPESWSCRKSTPRLVAWVS AAKVFIRDKLMERRNRTIGRT EKARIWEVTDRTVRTWIGEAV AAAAADGGFRVDLARRSIRK DRNARSQNPVHTEGDMNMNIK KIVKQATVLTFTTAFLAGGATQ AFAKENNQKAYKETYGVSHIT RHDMLQIPKQQQNEKYQVPQF DQSTIKNIESAKGLDVWDSWPL QNADGTVAEYNGYHVVFALA GSPKDADDTSIYMFYQKVGDN SIDISWKNAGRVF 4277 34645 B 4318 I 1374 4278 34646 A 4319 I 1293 4279 34647 A 4320 I 1278 4280 34648 A 4321 I 1254 MNMNIKKIVKQATVLTFTTALL AGGATQAFAKENNQKAYKETY GVSHITRHDMLQIPKQQQNEKY QVPQFDQSTIKNIESAKGLDVW DSWPLQNADGTVAEYNGYHV VFALAGSPKDADDTSIYMFYQ KVGDNSIDSWKNAGRVFKDSD KFDANDPILKDQTQEWSGSATF TSDGKIRLFYTDYSGKHYGKQS LTTAQVNVSKSDDTLKINGVN GYYCESVLFNKAIVYGGGTNFF RKESQKLQQSAKKRDAELANG ALGIIELNNDYTLKKVMKPLITS NTVTDEIERANVFKMNGKWYL FTDSRGSKMTIDGINSNDIYMG GYVSNSLTGPYKPLNKTGLVLQ MGLDPNDVTFTYSHFAVPQAK GNNVVITSYMTNRGFFEDKKA TFAPSFLMNIKGNKTSVVKNSIL EQGQLTVN 4281 34649 A 4322 I 726					ŀ		FFAPVFTMGKYYTQGDKVLMP
### AAKVFIRDKLMERRNRTGRT							LAIGGPLESPDRDGGPLESTNR
							DASPESWSCRKSTPRLVAWVS
AAAAADGGGFRVDLARRSIRK DRNARSQNPVHTEGDMNMNIK KIVKQATVLTFTTAFLAGGATV AFAKENNQKAYKETYGVSHIT RHDMLQIPKQQQNEKYQVPQF DQSTIKNIESAKGLDVWDSWPL QNADGTVAEVNGYHVVFALA GSPKDADDTSIYMFYQKVGDN SID\SWKNAGRVF							AAKVFIRDKLMERRNRRTGRT
DRNARSQNPVHTEGDMNMNIK KIVKQATVLTFTTAFLAGGATQ AFAKENNQKAYKETYGVSHIT RHDMLQIPKQQQNEKYQVPQF DQSTIKNIESAKGLDVWDSWPL QNADGTVAEYNGYHVVFALA GSPKDADDTSIYMFYQKVGDN SIDISWKNAGRVF 4277 34645 B 4318 1 1374 4278 34646 A 4319 1 1293 4279 34647 A 4320 I 1278 4280 34648 A 4321 I 1254 MNMNIKKIVKQATVLTFTTALL AGGATQAFAKENNQKAYKETY GVSHITRHDMLQIPKQQQNEKY QVPQFDQSTIKNIESAKGLDVW DSWPLQNADGTVAEYNGYHV VFALAGSPKDADDTSIYMFYQ KVGDNSIDSWKNAGRVFKDSD KFDANDPILKDQTQEWSGSATF TSDGKIRLFYTDYSGKHYGKQS LTTAQVNVSKDDTLKINGVN GYYCESSLFNKAIYYGGGTNFF RKESQKLQQSAKKRDAELANG ALGIIELNNDYTLKKVMKPLITS NTVTDEIERANVFKMNGK WYL FTDSRGSKMTIDGINSNDIYML GYVSNSLTGPYKPLNKTGLVLQ MGLDPNDVTFTYSHFAVPQAK GNNVVITSYMTNRGFFEDKKA TFAPSFLMNIKGNKTSVVKNSIL EQGQLTVN 4281 34649 A 4322 I 726							EKARIWEVTDRTVRTWIGEAV
KIVKQATVLTFTTAFLAGGATQ							AAAAADGGGFRVDLARRSIRK
AFAKENNQKAYKETYGVSHIT RHDMLQIPKQQQNEKYQVPQF DQSTIKNIESAKGLDVWDSWPL QNADGTVAEYNGYHVVFALA GSPKDADDTSIYMFYQKVGDN SIDISWKNAGRVF 4277 34645 B 4318 1 1293 4278 34646 A 4319 1 1293 4280 34647 A 4320 I 1278 4280 34648 A 4321 I 1254 MNMNIKKIVKQATVLTFITALL AGGATQAFAKENNQKAYKETY GVSHITRHDMLQIPKQQNEKY QVPQFDQSTIKNIESAKGLDVW DSWPLQNADGTVAEYNGYHV VFALAGSPKDADDTSIYMFYQ KVGDNSIDSWKNAGRVFKDSD KFPANDPILKDQTQEWSGSATF TSDGKIRLFYTDYSGKHYGKOS LTTAQVNVSKSDDTLKINGVN GYYCEES\LFNKA\YYGGGTNFF RKESQKLQQSAKKRDAELANG ALGIIELNNDYTLKKVMKPLITS NTVTDEIERANVFKMNGK WYL FTDSRGSKMTIDGINSNDIYML GYVSNSLTGPYKPLNKTGLVLQ MGLDPNDVTFTYSHFAVPQAK GNNVVITSYMTNRGFFEDKKA TFAPSFLMNIKGNKTSVVKNSIL EQGQLTVN 4281 34649 A 4322 I 726							DRNARSQNPVHTEGDMNMNIK
RHDMLQIPKQQNEKYQVPQF DQSTIKNIESAKGLDVWDSWPL QNADGTVAEYNGYHVVFALA GSPKDADDTSIYMFYQKVGDN SID\SWKNAGRVF 4277 34645 B 4318 I 1374 4278 34646 A 4319 I 1293 4279 34647 A 4320 I 1278 4280 34648 A 4321 I 1254 MNMNIKKIVKQATVLTFTTALL AGGATQAFAKENNQKAYKETY GVSHITRHDMLQIPKQQQNEKY QVPQFDQSTIKNIESAKGLDVW DSWPLQNADGTVAEYNGYHV VFALAGSPKDADDTSIYMFYQ KVGDNSIDSWKNAGRVFKDSD KFDANDPILKDQTQEWSGSATF TSDGKIRLFYTDYSGKHYGKQS LTTAQVNVSKSDDTLKINGVN GYYCEESLENKAIYYGGGTNFF RKESQKLQQSAKKRDAELANG ALGIIELNNDYTLKKVMKPLITS NTVTDEIERANVFKMNGKWYL FTDSRGSKMTIDGINSNDIYML GYVSNSLTGPYKPLNKTGLVLQ MGLDPNDVTFTYSHFAVPQAK GNNVVITSYMTNRGFFEDKKA TFAPSFLMNIKGNKTSVVKNSIL EQGQLTVN 4281 34649 A 4322 I 726							KIVKQATVLTFTTAFLAGGATQ
DQSTIKNIESAKGLDVWDSWPLQNADGTVAEYNGYHVVFALA GSPKDADDTSIYMFYQKVGDN SID\SWKNAGRVF 4277 34645 B 4318 1 1374 4278 34646 A 4319 1 1293 4279 34647 A 4320 I 1278 4280 34648 A 4321 I 1254 MNMNIKKIVKQATVLTFTTALL AGGATQAFAKENNQKAYKETY GVSHITRHDMLQIPKQQQNEKY QVPQFDQSTIKNIESAKGLDVW DSWPLQNADGTVAEYNGYHV VFALAGSPKDADDTSIYMFYQ KVGDNSIDSWKNAGRVFKDSD KFDANDPILKDQTQEWSGSATF TSDGKIRLFYTDYSGKHYGKQS LTTAQVNVSKSDDTLKINGV\N GYYCEES\LFNKA\YYGGGTNFF RKESQKLQQSAKKRDAELANG ALGIIELNNDYTLKKVMKPLITS NTVTDEIERANVFKMNGKWYL FTDSRGSKMTIDGINSNDIYML GYYSNSLTGPYKPLNKTGLVLQ MGLDPNDVTFTYSHFAVPQAK GNNVVITSYMTNRGFFEDKKA TFAPSFLMNIKGNKTSVVKNSIL EQGQLTVN 4281 34649 A 4322 I 726	İ		İ				AFAKENNQKAYKETYGVSHIT
QNADGTVAEYNGYHVVFALA GSPKDADDTSIYMFYQKVGDN SID\SWKNAGRVF	ļ						RHDMLQIPKQQQNEKYQVPQF
GSPKDADDTSIYMFYQKVGDN SID\SWKNAGRVF							DQSTIKNIESAKGLDVWDSWPL
SID\SWKNAGRVF							QNADGTVAEYNGYHVVFALA
1374 14278 34645 B 4318 1 1374 1293 14279 34647 A 4320 1 1278 1254 MNMNIKKIVKQATVLTFTTALL AGGATQAFAKENNQKAYKETY GVSHITRHDMLQIPKQQQNEKY QVPQFDQSTIKNIESAKGLDVW DSWPLQNADGTVAEYNGYHV VFALAGSPKDADDTSIYMFYQ KVGDNSIDSWKNAGRVFKDSD KFDANDPILKDQTQEWSGSATF TSDGKIRLFYTDYSGKHYGKQS LTTAQVNVSKSDDTLKINGVN GYYCEES\LFNKA\YYGGGTNFF RKESQKLQQSAKKDAELANG ALGIIELNNDYTLKKVMKPLITS NTVTDEIERANVFKMNGKWYL FTDSRGSKMTIDGINSNDIYML GYVSNSLTGPYKPLNKTGLVLQ MGLDPNDVTFTYSHFAVPQAK GNNVVITSYMTNRGFFEDKKA TFAPSFLMNIKGNKTSVVKNSIL EQGQLTVN 4281 34649 A 4322 1 726	İ						1
1							SID\SWKNAGRVF
1	4277	1			<u> </u>		
4280 34648 A 4321 I 1254 MNMNIKKIVKQATVLTFTTALL AGGATQAFAKENNQKAYKETY GVSHITRHDMLQIPKQQQNEKY QVPQFDQSTIKNIESAKGLDVW DSWPLQNADGTVAEYNGYHV VFALAGSPKDADDTSIYMFYQ KVGDNSIDSWKNAGRVFKDSD KFDANDPILKDQTQEWSGSATF TSDGKIRLFYTDYSGKHYGKQS LTTAQVNVSKSDDTLKINGV\N GYYCEES\LFNKA\YYGGGTNFF RKESQKLQQSAKKRDAELANG ALGIIELNNDYTLKKVMKPLITS NTVTDEIERANVFKMNGKWYL FTDSRGSKMTIDGINSNDIYML GYVSNSLTGPYKPLNKTGLVLQ MGLDPNDVTFTYSHFAVPQAK GNNVVITSYMTNRGFFEDKKA TFAPSFLMNIKGNKTSVVKNSIL EQGQLTVN			_				
AGGATQAFAKENNQKAYKETY GVSHITRHDMLQIPKQQQNEKY QVPQFDQSTIKNIESAKGLDVW DSWPLQNADGTVAEYNGYHV VFALAGSPKDADDTSIYMFYQ KVGDNSIDSWKNAGRVFKDSD KFDANDPILKDQTQEWSGSATF TSDGKIRLFYTDYSGKHYGKQS LTTAQVNVSKSDDTLKINGV\N GYYCEES\LFNKA\YYGGGTNFF RKESQKLQQSAKKRDAELANG ALGIIELNNDYTLKKVMKPLITS NTVTDEIERANVFKMNGKWYL FTDSRGSKMTIDGINSNDIYML GYVSNSLTGPYKPLNKTGLVLQ MGLDPNDVTFTYSHFAVPQAK GNNVVITSYMTNRGFFEDKKA TFAPSFLMNIKGNKTSVVKNSIL EQGQLTVN			—	<u> </u>	1		MNIMNIE E IVE O A TVI TETTA I I
GVSHITRHDMLQIPKQQQNEKY QVPQFDQSTIKNIESAKGLDVW DSWPLQNADGTVAEYNGYHV VFALAGSPKDADDTSIYMFYQ KVGDNSIDSWKNAGRVFKDSD KFDANDPILKDQTQEWSGSATF TSDGKIRLFYTDYSGKHYGKQS LTTAQVNVSKSDDTLKINGV\N GYYCEES\LFNKA\YYGGGTNFF RKESQKLQQSAKKRDAELANG ALGIIELNNDYTLKKVMKPLITS NTVTDEIERANVFKMNGKWYL FTDSRGSKMTIDGINSNDIYML GYVSNSLTGPYKPLNKTGLVLQ MGLDPNDVTFTYSHFAVPQAK GNNVVITSYMTNRGFFEDKKA TFAPSFLMNIKGNKTSVVKNSIL EQGQLTVN 4281 34649 A 4322 I 726	4280	34048	A	4321		1234	· · · · · · · · · · · · · · · · · · ·
QVPQFDQSTIKNIESAKGLDVW DSWPLQNADGTVAEYNGYHV VFALAGSPKDADDTSIYMFYQ KVGDNSIDSWKNAGRVFKDSD KFDANDPILKDQTQEWSGSATF TSDGKIRLFYTDYSGKHYGKQS LTTAQVNVSKSDDTLKINGV\N GYYCEES\LFNKA\YYGGGTNFF RKESQKLQQSAKKRDAELANG ALGIIELNNDYTLKKVMKPLITS NTVTDEIERANVFKMNGKWYL FTDSRGSKMTIDGINSNDIYML GYVSNSLTGPYKPLNKTGLVLQ MGLDPNDVTFTYSHFAVPQAK GNNVVITSYMTNRGFFEDKKA TFAPSFLMNIKGNKTSVVKNSIL EQGQLTVN 4281 34649 A 4322 1 726							, · · · · · · · · · · · · · · · · · · ·
DSWPLQNADGTVAEYNGYHV VFALAGSPKDADDTSIYMFYQ KVGDNSIDSWKNAGRVFKDSD KFDANDPILKDQTQEWSGSATF TSDGKIRLFYTDYSGKHYGKQS LTTAQVNVSKSDDTLKINGVIN GYYCEES\LFNKA\YYGGGTNFF RKESQKLQSAKKRDAELANG ALGIIELNNDYTLKKVMKPLITS NTVTDEIERANVFKMNGKWYL FTDSRGSKMTIDGINSNDIYML GYVSNSLTGPYKPLNKTGLVLQ MGLDPNDVTFTYSHFAVPQAK GNNVVITSYMTNRGFFEDKKA TFAPSFLMNIKGNKTSVVKNSIL EQGQLTVN 4281 34649 A 4322 I 726	ŀ						
VFALAGSPKDADDTSIYMFYQ KVGDNSIDSWKNAGRVFKDSD KFDANDPILKDQTQEWSGSATF TSDGKIRLFYTDYSGKHYGKQS LTTAQVNVSKSDDTLKINGV\N GYYCEES\LFNKA\YYGGGTNFF RKESQKLQQSAKKRDAELANG ALGIIELNNDYTLKKVMKPLITS NTVTDEIERANVFKMNGKWYL FTDSRGSKMTIDGINSNDIYML GYVSNSLTGPYKPLNKTGLVLQ MGLDPNDVTFTYSHFAVPQAK GNNVVITSYMTNRGFFEDKKA TFAPSFLMNIKGNKTSVVKNSIL EQGQLTVN 4281 34649 A 4322 I 726							[]
KVGDNSIDSWKNAGRVFKDSD KFDANDPILKDQTQEWSGSATF TSDGKIRLFYTDYSGKHYGKQS LTTAQVNVSKSDDTLKINGV\N GYYCEES\LFNKA\YYGGGTNFF RKESQKLQQSAKKRDAELANG ALGIIELNNDYTLKKVMKPLITS NTVTDEIERANVFKMNGKWYL FTDSRGSKMTIDGINSNDIYML GYVSNSLTGPYKPLNKTGLVLQ MGLDPNDVTFTYSHFAVPQAK GNNVVITSYMTNRGFFEDKKA TFAPSFLMNIKGNKTSVVKNSIL EQGQLTVN 4281 34649 A 4322 I 726							}
KFDANDPILKDQTQEWSGSATF TSDGKIRLFYTDYSGKHYGKQS LTTAQVNVSKSDDTLKINGVN GYYCEES\LFNKA\YYGGGTNFF RKESQKLQQSAKKRDAELANG ALGIIELNNDYTLKKVMKPLITS NTVTDEIERANVFKMNGKWYL FTDSRGSKMTIDGINSNDIYML GYVSNSLTGPYKPLNKTGLVLQ MGLDPNDVTFTYSHFAVPQAK GNNVVITSYMTNRGFFEDKKA TFAPSFLMNIKGNKTSVVKNSIL EQGQLTVN 4281 34649 A 4322 I 726							1
TSDGKIRLFYTDYSGKHYGKQS LTTAQVNVSKSDDTLKINGV\N GYYCES\LFNKA\YYGGGTNFF RKESQKLQQSAKKRDAELANG ALGIIELNNDYTLKKVMKPLITS NTVTDEIERANVFKMNGKWYL FTDSRGSKMTIDGINSNDIYML GYVSNSLTGPYKPLNKTGLVLQ MGLDPNDVTFTYSHFAVPQAK GNNVVITSYMTNRGFFEDKKA TFAPSFLMNIKGNKTSVVKNSIL EQGQLTVN 4281 34649 A 4322 I 726							ŀ
LTTAQVNVSKSDDTLKINGV\N GYYCEES\LFNKA\YYGGGTNFF RKESQKLQQSAKKRDAELANG ALGIIELNNDYTLKKVMKPLITS NTVTDEIERANVFKMNGKWYL FTDSRGSKMTIDGINSNDIYML GYVSNSLTGPYKPLNKTGLVLQ MGLDPNDVTFTYSHFAVPQAK GNNVVITSYMTNRGFFEDKKA TFAPSFLMNIKGNKTSVVKNSIL EQGQLTVN 4281 34649 A 4322 I 726	1						
GYYCES\LFNKA\YYGGGTNFF RKESQKLQQSAKKRDAELANG ALGIIELNNDYTLKKVMKPLITS NTVTDEIERANVFKMNGKWYL FTDSRGSKMTIDGINSNDIYML GYVSNSLTGPYKPLNKTGLVLQ MGLDPNDVTFTYSHFAVPQAK GNNVVITSYMTNRGFFEDKKA TFAPSFLMNIKGNKTSVVKNSIL EQGQLTVN 4281 34649 A 4322 I 726							1
RKESQKLQQSAKKRDAELANG ALGIIELNNDYTLKKVMKPLITS NTVTDEIERANVFKMNGKWYL FTDSRGSKMTIDGINSNDIYML GYVSNSLTGPYKPLNKTGLVLQ MGLDPNDVTFTYSHFAVPQAK GNNVVITSYMTNRGFFEDKKA TFAPSFLMNIKGNKTSVVKNSIL EQGQLTVN 4281 34649 A 4322 I 726					}		1
ALGIIELNNDYTLKKVMKPLITS NTVTDEIERANVFKMNGKWYL FTDSRGSKMTIDGINSNDIYML GYVSNSLTGPYKPLNKTGLVLQ MGLDPNDVTFTYSHFAVPQAK GNNVVITSYMTNRGFFEDKKA TFAPSFLMNIKGNKTSVVKNSIL EQGQLTVN 4281 34649 A 4322 I 726							1
NTVTDEIERANVFKMNGKWYL FTDSRGSKMTIDGINSNDIYML GYVSNSLTGPYKPLNKTGLVLQ MGLDPNDVTFTYSHFAVPQAK GNNVVITSYMTNRGFFEDKKA TFAPSFLMNIKGNKTSVVKNSIL EQGQLTVN 4281 34649 A 4322 I 726							1 7
FTDSRGSKMTIDGINSNDIYML GYVSNSLTGPYKPLNKTGLVLQ MGLDPNDVTFTYSHFAVPQAK GNNVVITSYMTNRGFFEDKKA TFAPSFLMNIKGNKTSVVKNSIL EQGQLTVN 4281 34649 A 4322 I 726							1
GYVSNSLTGPYKPLNKTGLVLQ MGLDPNDVTFTYSHFAVPQAK GNNVVITSYMTNRGFFEDKKA TFAPSFLMNIKGNKTSVVKNSIL EQGQLTVN 4281 34649 A 4322 I 726							· ·
MGLDPNDVTFTYSHFAVPQAK GNNVVITSYMTNRGFFEDKKA TFAPSFLMNIKGNKTSVVKNSIL EQGQLTVN 4281 34649 A 4322 I 726							1
GNNVVITSYMTNRGFFEDKKA TFAPSFLMNIKGNKTSVVKNSIL EQGQLTVN 4281 34649 A 4322 I 726							l *1
TFAPSFLMNIKGNKTSVVKNSIL EQGQLTVN 4281 34649 A 4322 I 726							`
4281 34649 A 4322 I FQGQLTVN							l '
4281 34649 A 4322 I 726							1
4282 34650 A 4323 1 1050	4281	34649	A	4322	1	726	
	4282	34650	Α	4323	1	1050	

SEQ ID	SEQ ID NO:	1	SEQ ID NO:	I.	1	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4283	34651	 A	4324	<u> </u> 	1185	 MNMNIKKIVKQATVLTFTTALL
4203	34031	^	1321		[1103	AGGATQAFAKENNQKAYKETY
						GVSHITRHDMLQIPKQQQNEKY
	1					
				•		QVPQFDQSTIKNIESAKGLDVW
					,	DSWPLQNADGTVAEYNGYHV
						VFALAGSPKDADDTSIYMFYQ
						KVGDNSIDSWKNAGRVFKDSD
						KFDANDPILKDQTQEWSGSATF
	1					TSDGKIRLFYTDYSGKHYGKQS
	1	ŀ				LTTAQVNVSKSDDTLKINGVED
		ł				HKTIFDGDGKTYQNVQQFIDEG
						NYTSGDNHTLRDPHYVEDKGH
						KYLVFEANTGTENGYQGEESLF
Ì						NKAYYGGGTNFFRKESQKLQQ
						SAKKRDAELANGALGIIELNND
					,	YT\LKKVMKPLITSN/TVPQAKG
		1				NNVVITSYMTNRGFFEDKKATF
		l				APSFLMNIKGNKTSVVKNSILE
						QGQLTVN
4284	34652	В	4325	1	867	
4285	34653	Α	4326	1	495	
4286	34654	Α	4327	3	1394	GDMNMNIKKIVKQATVLTFTT
						A/LLAGGATQAFAKENNQKAY
						KETYGVSHITRHDMLQIPKQQQ
						NEKYQVPQFDQSTIKNIESAKG
						LDVWDSWPLQNADGTVAEYN
	•	1				GYHVVFALAGSPKDADDTSIY
		İ		ŀ		MFYQKVGDNSIDSWKNAGRVF
		ŀ		•		KDSDKFDANDPILKDQTQEWS
						GSATFTSDGKIRLFYTDYSGKH
						YGKQSLTTAQVNVSKSDDTLKI
		ŀ				NGVEDHKTIFDGDGKTYQNVQ
						QFIDEGNYTGDPLEAETAVINH
		1				KKRKNSPRIVQSNDLTEAAYSL
	1			1		SRDQKRMLYLFVDQIRKSDGTL
						QEHDGICEIHVAKYAEIFGLTSA
	,					EASKDIRQALKSFAGKEVVFYR
		1				PEEDAGDEKGYESFPWFIKRAH
		1]			SPSRGLYSVHINPYLIPFFIGLQN
		1				RFTQFRLSETKEITNPYAMRLY
						ESLCQYRKPDGSGIVSLKIDWII
						ERYQLPQSYQRTPDFRRRFLQV
				<u></u>		CVNEING
4287	34655	В	4328	9	1004	
4288	34656	Α	4329	1	768	
4289	34657	A	4330	1	1308	
4290	34658	В	4331	58	753	
4291	34659	В	4332	1	409	
4292	34660	В	4333	1	921	
4293	34661	A	4334	1	1026	
4294	34662	В	4335	1	945	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4295	34663	A	4336	11	528	MNMNIKKIVKQATVLTFTTALL
1293	3 1003	` `	1330	!		AGGATQAFAKENNQKAYKETY
						GVSHITRHDMLQIPKQQQNEKY
		İ				QVPQFDQSTIKNIESAKGLDVW
						DSWPLQNADGTVAEYNGYHV
		ŀ				VFALAGSPKDADDTSIYMFY/Q
						KDQTQEWS\GSATFTSDGKIRLF
		1				YTDYSGKHYGKQSLDTA\Q*NV
						VKSG
4296	34664	Α	4337	l	1701	
4297	34665	В	4338	97	1449	
4298	34666	Α	4339	1	1581	
4299	34667	В	4340	1	1539	
4300	34668	Α	4341	87	1078	SLPNLDNAAICSSSSSPTRTR*SL
						SEGATQ\AFAKEKYPHKHTKKR
						SGVFHITRHDMLQIPKQQQNEK
		•				YQVPQFDQSTIKNIESAKALDV
		1				WDSWPLQNADGTVAEYNGYH
						VVFALAGSPKDADDTSIYMFY
				,		QKVGDNSIDSWKNAGRVFKDS
						DKFDANDPILKDQTQEWSGSA
		ŀ		ŀ		TFTSDGKIRLFYTDYSGKHYGK
				•		QSLTTAQVNVSKSDDTLKINGV
						EDHKTIFDGDGKTYQNVQQFID EGNYTSGDNHTLRDPHYVEDK
						GHKYRGPLESPSTHQAEFNPTS
						CVSSLGTLQGFPAPAWLALAHP
						VHPLKHKSGGSNRLSAAIWGIK
						RKPAR
4301	34669	A	4342	1	1344	
4302	34670	Α	4343	1	1713	
4303	34671	Α	4344	3	1918	
4304	34672	A	4345	254	1118	RPPAFAKK*PKAYKET/YGVSHI
						TRHDMLQIPKQQQNEKYQVPQ
						FDQSTIKNIESAKGLDVWDSWP
						LQNADGTVAEYNGYHVVFALA
						GSPKDADDTSIYMFYQKVGDN
•						SIDSWKNAGRVFKDSDKFDAN
						DPILKDQTQEWSGSATFTSDGK
						IRLFYTDYSGKHYGKQSLTTAQ
						VNVSKSDDTLKINGVEDHKTIF
1						DGDGKTYQNVQQFIDEGNYTS
						GDNHTLRDPHYVGGTSWEPGV
1						FSVSCVFFGQQEGV/HG*DEFLD
						FSYWFQGG*ICLYQKAS*QNTT
L						SYKRYTGS

SEQ ID		•	SEQ ID NO:		I .	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	sequence		07/510,217	sequence	or pepulae soquesie	ection, v possou serveside insertion,
4305	34673	A	4346	1	1952	MNMNIKKIVKQATVLTFTTALL
						AGGATQAFAKENNQKAYKETY
	İ					GVSHITRHDMLQIPKQQQNEKY
						QVPQFDQSTIKNIESAKGLDVW
ļ						DSWPLQNADGTVAEYNGYHV
			Ì			VFALAGSPKDADDTSIYMFYQ
						KVGDNSIDSWKNAGRVFKDSD
						KFDANDPILKDQTQEWSGSATF
						TSDGKIRLFYTGSLNSSKTEKY
				:		QVPHIDQSTIKNIESAKGLDVW
						DSWPLQNADGTVAEYNGYHV
	1					VFALAGSPKDADDTSIYMFYQ
						KVGDNSIDSWKNAGRVFKDSD
						KFDANDPILKDQTQEWSGSATF
						TSDGKIRLFYTDYSGKHYGKQS
						LTTAQVNVSKSDDTLKINGVED
<u> </u>						HKTIFDGDGKTYQNVQQFIDEG
Ì					·	NYTSGDNHTLRDPHYVEDKGH
						KYLVFQDHTGTEEHPQPQ\ERP
						RTQSFTSAFAERRECIPNVPADT
						KLSKIKTLRLATSYIAYLMDLL
						AKDDQNGEAEAFKAEIKKTDV
						KEEKRKKELASKCLDLEQLGAS
						VEPTGNLRTKITKEKPRHTGPPE
						VVVPGCCPHRSRAYKSDKYAH
						TLTVTASQHAPPPPTHMEGFEL
						FHLPDLCSPSQDAQTTGRTQMK
		ļ.				PDHSPRPSHRVPQAKGNNVVIT
Ì						SYMTNRGFFEDKKATFAPSFLM
		<u></u>				NIKGNKTSVVKNSILEQGQLTV
4306	34674	Α	4347	1	1029	
4307	34675	Α	4348	276	1248	CVWLGCRGYYPKAYKETY\GV
						SHITRHDMLQIPKQQQNEKYQV
:						PQFDQSTIKNIESAKGLDVWDS
ŀ						WPLQNADGTVAEYNGYHVVF
						ALAGSPKDADDTSIYMFYQKV
						GDNSIDSWKNAGRVFKDSDKF
						DANDPILKDQTQEWSGSATFTS
						DGKIRLFYTDYSGKHYGKQSLT
						TAQQLLQLVQFQEVDTDFDFPE
						EDKKEEFEECLEKFFSTGPARPP
						TKEKVKRRVLIEPGMPLNHIEY
						CNHEIMGKNVYYKHRWVAEH
						YFLLMQYDELQKICYNEFVPSV
!						IFLRYKSPGEAAGTCHLKQRRW
						VMPEAAAPVGTGSRYPLTGQL
4308	34676	A	4349	1	242	MNSIQIPKQQQNEKYQVPQFDQ
						STIKYIESPKELDVWDSWPLQN
						ADGTVAEYNGYHVAFALAG/S
						PKDADDTSIYMFYQKI
4309	34677	В	4350	1	2198	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4310	34678	A	4351	1	2796	
4311	34679	Α	4352	2047	3531	
4312	34680	A	4353	1	3336	
4313	34681	Α	4354	1	1409	MKRAPVIPKHTLNTQPVEDTSL
						STPAAPMVDSLIARVGVMARG
						NAITLPVCGRDVKFTLEVLRGD
						SVEKTSRVWSGNERDQELLTE
	ŀ			, ·		DALDDLIPSFLLTGQQTPAFGR
						RVSGVIEIADGSRRRKAAALTE
						SDYRVLVGELDDEQMAALSRL
						GGATQAFAKENNQK\AYKETY
						GVSHITRHDMLQIPKQQQNEKY
						QVPQFDQSTIKNIESAKGLDVW
1		1				DSWPLQNADGTVAEYNGYHV
						VFALAGSPKDADDTSIYMFYQ
						KVGDNSIDSWKNAGRVFKDSD
	1					KFDANDPILKDQTQEWSGSATF
						TSDGKIRLFYTDYSGKHYGKQS
						LTTAQVNVSKSDDTLKINGVED
						HKTIFDGDGKTYQNVQQFIDGY
						LLEPDGGALQNFQRYTGIQHVH
						RIGMAERMWCDRNRERHTVSS
						SGGNRLPNPGPDRSVRHFPDPR
						FLCPSCATVTPLHELIANKYLSG
						KIGAKKLRKLLIKHVD
4314	34682	Α	4355	1	2316	
4315	34683	Α	4356	93	924	AQTDAAEKSVSIAQLFQACLSIF
						SSGDV/AGGATQAFAKENNQK
						AYKETYGVSHITRHDMLQIPKQ
						QQNEKYQVPQFDQSTIKNIESA
						KGLDVWDSWPLQNADGTVAE
						YNGYHVVFALAGSPKDADDTS
	1	1]	IYMFYQKVGDNSIDSWKNAGR
	1					VFKDSDKFDANDPILKDQTQE
	1			•		WSGSATFTSDGKIRLFYTDYSG
						KHYGKQSLTTAQVNVSKSNDT
						LKINGVGKYKTIFDGDGKTYQT
						VQQFIDEGNYTSGGHHTL\KDP
]			SYNPPLDLSGGNSGYQSQET

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO: in USSN	Nucleotide location of first		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4316	34684	A	4357	1	3118	MNMNIKKIVKQATVLTFTTALL
	}					AGGATQAFAKENNQKAYKETY
						GVSHITRHDMLQIPKQQQNEKY
		l				QVPQFDQSTIKNIESAKGLDVW
		ŀ				DSWPLQNADGTVAEYNGYHV
						VFALAGSPKDADDTSIYMFYQ
						KVGDNSIDSWKNAGRVFKDSD
						KFDANDPILKDQTQEWSGSATF
						TSDGKIRLFYTDYSGKHYGKQS
1		ł				LTTAQVNVSKSDDTLKINGVED
				ļ		HKTIFDGDGKTYQNVQQFIDEG
]		NYTSGDNHTL\RDP\HYVENKG
						HKYLGFETNTGTENGYQGEESL
					ļ	FNKAYYGGGTNFFRKESQKLQ
]		l				QSAKKRDAELANGALGIIELNN
]						DYTLKKVMKPLITSNTVTDEIE
						RANVFKMNGKWYLFTDSRGSK
		ļ				MTIDGINSNDIYMLGSDESPND
						FGNRHLHKERLAVYRWHASFI
						CSGNTMPIVLVDWSDIREQKRL
		ľ				MVLRASVALHGRSVTLYEKAF
						PLSEQCSKKAHDQFLADLASIL
						PSNTTPLIVSDAGFKVPWYKSV
				·		EKLGWYWLSRVRGKVQYADL
						GAENWKPISNLHDMSSSHSKTL
						GYKRLTKSNPISCQILLYKSRSK
						GRKNQRSTRTHCHHPSPKIYSA SAKEPWVLATNLPVEIRTPKQL
		ŀ		,		VNIYSKRMQIEETFRDLKSPAY
						GLGLRHSRTSSSERFDIMLLIAL
						MLQLTCWLAGVHAQKQ
4317	34685	Α	4358	1	1326	
4318	34686	Α	4359	2140	4390	
4319	34687	В	4360	1	7271	
4320	34688	Α	4361	1	1729	
4321	34689	Α	4362	5118	5687	
4322	34690	В	4363	1	4726	
4323	34691	В	4364	1	3688	
4324	34692	Α	4365	1	1401	
4325	34693	Α	4366	1	1932	
4326	34694	Α	4367	1	1407	
4327	34695	Λ	4368	1	1491	
4328	34696	В	4369	1	855	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hođ	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	į	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	:			sequence		
4329	34697	Α	4370	137	1014	ASEGKQMLRDFVTTRSALKELL
						KRALNMARNNQYQPLQKHAK
					1	L*RPSML*RNCIN*QPGRDT/TN
						KKENFRPISLMNIDAKILNKILA
						NRIQQHIKKLIHNDQVGFIPGM
						QGWFNIHKSINIIHHIKRTNDKN
						HMIISIDAEKAFNKIQQPFMLKT
				•		LNKLGIDGTYLKIIRAIYDRPTA
				·	}	NVILNGQKLEAFPFKTGTRQGC
						PLSPLLFNIVLKVLARAIRQETEI
						KGIQLAKEEVKLSLFADDMIVY
						LENPIISAQNLLKLISNFSK/VSG
						YKINVQKSQAFLYTINRQTESQI
4330	34698	Α	4371	3	1234	
4331	34699	Α	4372	1	2850	MGMGPAKPGMGGNLLVCWLQ
						RPWEKRSIWAEVYRSSRYSHS
						WLPLSRKGCDFSGTCRQTLSIL
						TQPLRQWGLEGIKKPNSWIISEE
						SVSNGGPPLLIPRQTASGVDLQ
						QTPTDLQLRVLTVRRKTNKQK
						GIASTSTKRTSTPKPHLYVTIIK
	ļ					DQSYIKPQRWGKNIAEKLKILKI
						RVALSLQRNAAPHQQWNKAG
		ļ				RRMSLMSSQKKASEVIESQMN
						EIKGEEKFREKRVKRNEQSLQEI
						WDYVKRPDLRLIGVPD
4332	34700	В	4373	16	701	
4333	34701	В	4374	227	3743	WANT A EVENTA OF ELECT DECC
4334	34702	Α	4375	221	686 .	KVMLAEYPVFAQLTLTLPPSSA
						SWEPSRGPGPRGIRGSCPEWLA
						SGPG\KAAPGAGVPPPAASFPDP
						PPRLRAPALAVSRGLRRELPSG
					-	LDWTHCLRTLPSLIVQILQQAA
						LLGLPPAYSDQLQRAGQLHFYS
1225	0.4800	ļ.,	1076	016		GLIKISLVLTTRLSFWGTTE
4335	34703	Α	4376	216	644	VTYSKEKECGEVADSVÄKTAL
						EKDGAPRTGDPRPNLGADPPRS
						LVSSAGPQAVRPVKPARQFPPQ
						PPRYSQGPARAAGEEGRGMRPP
						GAGRRLPGPPLPGPEASHSGQL
					,	PLM/PPGPGPRLGSQEPVSLSRY
		<u> </u>		<u> </u>		LQTQARMPGPRP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				<u> </u>		
4336	34704	Α	4377	18	1023	QIQHSPLVSPLPSLPPQPLVAEE
						EPPVA/PWPRRLLPATSTSH/PSH
						PLTEPVPPT\SGRGCCLWTKRQ
						QKMCRTTYTSVGRK/CTFPIDS
l						GALRLSDGEMRALQTPTGPQST
						VEGHTHLQSL/PHHDRVTATPG
						TEPGLRAAGNRRIFYPGP/VTSQ
		1				VQPQLLCGYGNASRTPAALTPG
						PAPPTQASLPNCGICPHLQMGR
İ				ŀ		PTSPC/PPEGGHPSSSLYISLSPPP
						PSAPALRLPPPLP\SAPTAPAL/P/
						PAAPSAPALRLLRPLPCCSFRPR
						PAAPSAPAPRLLPPLLVCSFRPC
						PAAPSAPAGLLLPPLLCCSGLSP
	i					RLCCPHSSCSSDPPRLQRKADSS
4337	34705	В	4378	1	984	
4338	34706	A	4379	332	847	VKLLLQDKEICILCQKTVYPME
						CLVADKQNFHKSCF\RCHHCNS
						KLSLGNYASLHGQIYCKPHFKP
						TFQNPKGNYDEGFGHKA\HKD
=						RWNWQKPKADSVDFIPNEEPN
						MCKNIAENTLVPGDRNEHLDA
						GNSEGQRNDLRKLGERGKLKV
İ						IWPPSKEIPKKTLPFEEELKMSK
4339	34707	Α	4380	305	505	GNLERMLNLGMVKQQKLPAIM
Ì						KTQVLML*AINVPAKPLFPQSG
						GAVRTTHGGKSRLKETGATSD
						TE
4340	34708	Α	4381	56	260	IVKTQSIDG/MGNLRITEKGLKL
	1					EGDS/EFLQPLYAKEIQSRPGLG
						TQEQSCQTLSSCSSRGQQQHAE
4341	34709	Α	4382	137	920	
4342	34710	Α	4383	532	1680	LLTTRTSFRSENHRHVGLLLVM
						TDNTRDKEYFGDESKRENEEKT
					-	VEKSIGEKQATLTTHANIITIRH
						CVKPEPDFSDHLNLLLGRADIT
		İ]		GEEMAAQRSSVEKLANGNIAL
						VDSLRSRSLEEGDSDPHKRLSG
						AQDIKTTVEEVIADVVEIARELE
						LEVEPEDVTEFLQAHEKTLTDV
						ELFLINEQIKWFLEMKSTPRED
						AVIIAETITKVLEYDINLVTKQQ
						QGMRQLTPILKEVLLWVKCHQ
				1	ļ	TALHATEKPFIKGRINPCGKIHT
						CLNLRNCGQLLIREEEEEDKEE
						EEQYEEKEEEEEEEEEEEE
		[}		EEEEEEEEEEEEEEEEEEEEEEEEEEEEEEEEE
						DEEEDKEKEEEEDNKEEEEEED
						KEKEEEEDKE\EKEDKEEEEDK
	<u> </u>	1	L	L		REALIZED RECEIVED REFERENCE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	•	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence	į	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	1			sequence		
4343	34711	A	4384	3	495	EDTGTFRIY\ESAGAVKKARGF
						LEFVEDFIQVSKNLIGKVIGKNG
		[KVIQEIVDKSDMVPVRIEGDSE
	i					NKLPREDKDDRDSRHQRDSRR
						CPGGRCRSVSGRRGRGGPRGG
						KSSISSVPKDPDSNPYSVLDN/T
						ESDQTADTDASKSHHSTNRHTR
•		ŀ				SRRRTDEDAVL
4344	34712	Α	4385	ì	550	TESERKDELSDWSLAGEDDRDS
						RHQRDSRRRPGGRGRSVSGGR
				:		GRGGPRGGKSSISSVLKDPDSN
						PYSLLDNTESDQTADTDASESH
						HSTNRRRRSRRRRTDEDAVLM
				,		DGMT\ESDTASVNENGL\AKDV
!						IEEHGPSEKAINGPT\SASG\DDIS
1						KLQRTPGERKRLIP*KKENTQE
						AAVLNGVS
4345	34713	Α	4386	1	2063	MAELTVEVRGSNGAFYKGFIK
						DVHEDSLTVVFENNWQPERQV
						PFNEVRLPPPPDIKKEISEGDEV
						EVYSRANDQEPCGWWLAKVR
						MMKGEFYVIEYAACDATYNEI
1		ŀ				VTFERLRPVNQNKTVKKNTFFK
						CTVDVPEDLREACANENAHKD
	1					FKKAVGACRIFYHPETTQLMIL
						SASEATVKRVNILSDMHLRSIR
				·		TKLMLMSRNEEATKHLECTKQ
						LAAAFHEEFVVREDLMGLAIGT
						HGSNIQQARKVPGVTAIELDED
	1					TGTFRIYGESADAVKKARGFLE
		·				FVEDFIQVPRNLVGKVIGKNGK
						VIQEIVDKSGVVRVRIEGDNEN
				E - -		KLPREDGMVPFVFVGTKESIGN
						VQVLLEYHIAYLKEVEQLRME
						RLQIDEQLRQIGMGFRPSSTRGP
						EKEKGYATDESTVSSVQGSRSY
						SGRGRGRRGPNYTSGYGTNSEL
		ŀ				SNPSETESERKDELSDWSLAGE
					<u>.</u>	DDRDSRHQRDSRRRPGGRGRS
				ŀ		VSGGRGRGGPRGGKSSISSVQY
						RSNIHNCSTLKRIFLASDMNIVL
						KDPDSNPYSLLDNTESDQTADT
	1					DASESHHSTNRRRRSRRRRTDE
						DAVLMDGMTESDTASVNENGL
			<u> </u>			DDSEKKPQRRNRSRRRRFRGQ
						AE\DRQPAIDFIYKEVEKVVSL
	1					WQAKDVIEEHGPSEKAINGPTS
	1	L				ASGDDISKLQRTPGEEKINTLKE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
4346	34714	Α	4387]	1882	CGSNMADVTVEVRGSNGAFYK
						GFIKDVHEDSLTVVFENNWQPE
						RQVPFNEVRLPPPPDIKKEISEG
						DEVEVYSRANDQEPCGWWLA
						KVRMMKGEFYVIEYAACDATY
						NEIVTFERLRPVNQNKTVKKNT
]		1				FFKCTVDVPEDLREACANENA
						HKDFKKAVGACRIFYHPETTQL
						MILSASEATVKRVNILSDMHLR
						SIRTKLMLMSRNEEATKHLECT
						KQLAAAFHEEFVVREDLMGLA
						IGTHGSNIQQARKVPGVTAIEL
						DEDTGTFRIYGESADAVKKAR
						GFLEFVEDFIQVPRNLVGKVIG
						KNGKVIQEIVDKSGVVRVRIEG
						DNENKLPREDGMVPFVFVGTK
						ESIGNVQVLLEYHIAYLKEVEQ
						LRMERLQIDEQLRQIGSRSYSG
						RGRGRRGPNYTSGYGTNSELSN
						PSETESERKDELSDWSLAGEDN
						RDSRHQRDSRRRPGGRGRSVSG
						GRGRGGPR\GGKSSISSVLKDPD
						SNPYSLLDNT\ESDQTADTDASE
						SHHSTNRRRR/SIRRRRTD\EDA
		1		•		VLMNGMTESDTASVNENGLVT
						VADYISRAESQSRQRNLPRETL
						AKNKKEMAKDVIEEHGPSEKAI
						NGPTSASGDDISKLQRTPGEEKI
						NTLKEENTQEAAVLNGVS
4347	34715	Α	4388	2	421	PRVRDSDTEDDSEAEHFESFIHP
						TAMMFTSTINLLQTLCLSAGVH
						AEIMQSEATKTLCGLL\KSSPNR
						LVYREQHRSWCTLGFVQSIALT
						LQVCGALSSLQWITLLMKVVE
						GHAPFTATSLQRQILAVHLLQA
						VLPSWDK
4348	34716	Α	4389	269	417	DLNCKVGSCFEVYSS*KQGIN*I
						KLGDSKT*P*LSGPTSENLKNSS
						LAE
4349	34717	Α	4390	1	516	
4350	34718	С	4391	1	1527	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
İ	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
4351	34719	IA	4392	200	1267	TFSKASRGGNPHSMTKAPSDFR
7551	34717		14372	200	1207	KARQTGIPGCSQLGSRYSLEPE
						QSALRLVCIQKLQESSTTCEDFF
						CPLCGRAWAVSTPLTDSPSPGH
						QPAVK*LGLVPFSDTHHPLPFQ
						VLSTDDTSSSSSCSSSCSASSSSP
						/SLLLLLFLLLLLLLLMLLLKL
						FLLLLLFLL/RPPASPPLLPPALS
						PPL/HCSSSPSAPPASPPPAPPPPP
						APPPSPPPAPPPSAPSSPAPLPPA
						PASPPFSCSSSSCSSSCSSSFSSC
						SSLSSSAQAEGSLRAPRESSPSL
						DPSAPQRVKVVPPQAGSGHRA
		ĺ				GGALENRPRGKKPWLHFRPGL
ŀ						RSRLPARSLRSRPAPTRWRLRSS
						GRFTGAATATATATT
4352	34720	Α	4393	1	2607	MMGHSSAIPLTATPGELKGQSP
1332	3 1,20	ļ.,	.5,5	<u> </u>	2007	TKMPDPELGCQGAKSQGCSRN
						ARHQKARSMPLQDQHLALAIL
						LELAVQRGTLSQMLSAILLLLQ
						LWDSRAQETDNERSAQGTSTL
1						LLSLLQTFQSIICSKDTPPSEGN
						MHLLSGPLSPSESFLRESFFTVQ
		-				NCRNNEEVTLICKADLENHNK
						DGGFWIVIDEKVYDIKDFQTQS
1						LTGNSILAQFAGENPVVALEAA
1						FEFEVTRESMHAFCVGQYLEVR
						LYALSDAEDGRG\TL*WLQSSIF
						SG/GLQTSQIHYSYNEEKDEDH
						CS/SPVGTPASKSR\CSHRWALG
						DHSQAFLQAIADNNIQDHNVKT
						HQEQGRSYKEVCTPVIERLRFL
						SNELRPAVGNDLSIISEFKLLSSL
						PRWRRIAQKIIRERRKKRIPKKP
						ESTADEEKIGNEESDLEEACILP
						HSPINVDKRPIAIKSPKTITSENP
						LGPSLGSIPQARFLLMMLSMLT
						LQHSANNLDLLLNSGTLALTQT
						ALRLIGPSCDNVEEDMNASAQ
			}			GVSATVLEATRKETAPVHLPVS
				ŀ		GPELAATMKIGTRVMRGVDW
						KWGDQDGPPPGLGRVIGELGE
			1			DGWIRVQWGTGSTNSYRMGK
						EGKYDLKLAELPAAAQPSAEDS
						DTEDDSASPNRLVYREQHRSW
			1			CMLGFVRSIALTPQVCGALSSP
						QWITLLMKVMKGHAPFPAASL
						QRQRWVAVSLPHALVKSGTVP
	L		<u> </u>	<u> </u>		ZIZIKW VA USETTAL VRSGTVF

	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	1	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
4353	34721	Α	4394	266	1110	WARGGCARNALASGNAIQGGK
						CNPGLFPPSPNRLVYREQHRSW
						CMLGFVRSIALTPQVCGALSSP
						QWITLLMKVMKGHAPFPAASL
						QRQLCPE/HTSCPVLKDFCKSVI
	ŀ					TDVACSSLISTLLVFWGGLLHT
ļ		1				HKEASESWREAKSTSYVAAAR
					:	ENEEDAKAEPPTPGIKPSDLVRL
						IHYQENSMGETAPMIQIISHWV
						PPTTHGIYGSTIQDEIRVGVSYP
						GHTDARGFQLLLVSGDFSIPYW
		•				SLSSAYTSVNSSFVESLQSNLLK
						GILLPATIMTDPRTTGHQ
4354	34722	Α	4395	1	734	MVQLSGKRILNSPYLELRCHQN
		1				MDHLGWVIKKSLNRSEVSWVP
ļ						GLEFPWGPKPFREVIAGPLLRN
						NGQSLESSSLEGSHVGVYFSAH
İ						WCPPCRSLTRVLVESYRKIKEA
			į			GQNF\EIIFVSADRSEESFKQYFS
						EMPWLAVPYTDEARRSRLNRL
						YGIQAHLFLTANAEDFDTTVQV
		İ				NKIILITYRQNENSLSSLKTGET
		į				EAQGRLQGSPSNVRGHDPDRH
						AIPLSVNRWNPSKSSPSPAVWS
4355	34723	Λ	4396	195	1071	LHEFDSSRDLTSGLGGARTHRR
		İ				LGGPSDAPRGLPAPPPAPPVRPG
						/PRSPGPSAGTAR/DAPRPSVQM
						RAQRPARGSTKDLIETCCAAGQ
						QWAIDNDECLEIPESGTEDNVC
		:				RTAQRHCCVSYLQEKSCMAGV
						LGAKEGETCGAEDNDSCGISLY
						KASLTCGLQGRCLNPQQASMG
		İ				LFSYDVQSSKKINRSIQEKLGG
						HGVCAATPGGGMRNCGRLRRS
	·					GQRRGGTDRCEAVLTGLFTRA
						LIREQMGDPHPLDHTGQLAKPL
						EVEKTPARWKYLDTNGEKEEP
						ELRTQCPSLYED
4356	34724	Α	4397	1	520	MMGEKAEKPDTKEKKPKAKK
						ADAGGKRNCRYSRSAMYSRKT
						TSRKKYSAAKSKVEKKKKFLA
						TVTKPVGVDKNSGTQVVKLHK
		1				MPRYYPTEDVPLKLLSHG/KKK
						PFSQHR/RRVVFLKQLV/SGTGP
						LVLNQVPLRRTHQKFVIATSTKI
						GSSNVKIAKRLTGAYFKKVWK
					_	PKHQE
	34725	-	4398	67	243	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4358	34726	A	4399	1	333	QRSCIENILRAC/VGLPPQNHML
						LEHKM\DAKRVGPVAATYPML
						NKKGPVPAATNGCTGDANGHL
						QEEPPMPTT*GPGHTVSRLFLPA
		ļ				APHDPTLKAPTNNSAATQPSKN
						KKK
4359	34727	Α	4400	587	1013	GAASAGRGPGPRAPGLWGRGP
		ĺ				AAAGASLVPTDHVHLSYNHLG
						NNDGENLSAP/SQFRSKEVSKS
						NVVDD/MVQSNPVLYTPGEEPD
			,			HATRCWPHPSAGPSAADRAVP
	1					ARPAGAPATEPHAPGTQNGAP
						GPSLKRVGPVAATYPI
4360	34728	Α	4401	2	334	
4361	34729	В	4402	257	975	
4362	34730	Α	4403	30	365	EEAETVLVGQLKQLSSCLAVH
						KYRPETKQEKKQRLLARAEKK
						AAGKGDVPTKRPPVLRAGVNT
						VTTLVENKKAQLV\CRKMGVP
						YCIIKGKARLGRLVHRKTCTTV
		<u> </u>				AFTQVN
4363	34731	C	4404	62	217	
4364	34732	Α	4405	2	69	
4365	34733	Α	4406	1	951	GTRPKMPKGKKAKGKKVAPAP
						AVVKKQEGFRKKW*IPWFEKR
						P\KNFGIGQDIQPQKRPPPLL*K
		ŀ				WPRQYQACSGQRAILYKR\LKV
						PPAMKPVSPRALD\RQT\ATQLA
		ŀ				*AVAHKVQTQRQKQEKKQRL\
					+	LARADEEGCLAKGDVPNERDP
					•	PVPSSQEFNPVSPPLVKEQEKLK
						LVVNWHTDV\DPHPSLVCLPC/
						LCPAPVS*KMGGPFTCIIQGKRA
						RLWDRLVPQERPCTTCPPFT\QV
						N\SEDKVRLLAKAGLEAIQGPIY
						N*PDTMEIRPSLGVGNVLG\PKS
						VARI\AKARNRHKAKETATHTG
			1			LNVTLLSFLYYKNN

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4366	34734	Α	4407	1	1392	MPDVSEEQKESVCTGSMMREE
'500	.,,,,,	ļ.,			,	ESSRKGKVRTAGAKSSSSDRVP
ļ						RLNQEEVESLNRPITGAEIVAIIN
						SLPTKKSPAPDGFTAEFYQRIRI
		ļ				QQPIIRIQQPIKKLIQHDKVGFIP
					İ	GMQGWFNICKSINVIQHINRTK
						EKNHMIISIDAEKPFDKIQQCFM
						LKTLNKLGIDGTYLKIIRAIYHK
						PTANIILNGQKLEAFPLKTGTRQ
ļ			:			GCPLSPLLFNIVLEVLARAIRQE
						KEIKGIQLGKEEVKLSLFADDM
						IIYLENPTVSAQNLLKLISNFSK
						VSGYKINVQKSQAFLYTNNRQ
						TESQIMSELPFTIASKRIKHLGIQ
						LTRDVKDLFKENYKPLLNEIKE
						DTKKWKNIPCSWAGRISIMKM
						AILPKVIYRFNAIPIKLPMTFFTE
						LEKTTLKFIRNQKRAHIAKSILS
						QKNKAGGITLPDFKLYYKATV
						TKTAWYWYQNRDIDQCTRTQP
						\SEITPHIYNYLIF
4367	34735	Α	4408	1	1947	MALRRLSHDVSGALLLANGES
						TGNSGGSSGSSPSGGATSGSSQ
						TSISGDVVEACCSVLSMVCADP
						VYKVYVAAL\QCMLLVTLEDPS
						SHFTRMRRRLM/AYADEVEIAE
						AIQLGVEDTLDGQQDSF\CRHL
						FPTTIWKPQRTVP/LECTIHLEKT
						GKGLCATKLSASSEDISERLASI
						SVGPSSSTTTTTTTEQPKPMVQ
						TKGRPHSQCLNSSPLSHHSQLM
						FPALSTPSSSTPSVPAGTATDVS KHRLQGFIPCRIPSASPQTQRKF
						SLQFHRNCPENKDSDKLSPVFT
				ł		QSRPLPSSNIHRPKPSRPTPGNTS
				j		KQGDPSKNSMTLDLNSSSKCD
						DSFGCSSNSS/NCCYT\SDETVFT
						PVEEKCRLDVNTELNSSIEDLLE
		İ				ASMPSSDTTVTFKSEVAVLSPE
		1				KAENDDTYKDDVNHNQKCKE
ł		ĺ				KMEAEEEEALAIAMAMSASQD
		l				ALPIVPQLQVENGEDIIIIQQDM
					}	TFFRHIIPPIQWIYKKESANLLID
	}					STGQRLRIADFGAAARLASKGT
						GAGEFQGQLLGTIAFMAPEVLR
						GQQYGRSCDVWSVGCAIIEMA
						CAKPPWNAEKHSNHLALIFKKL
						LDFANTACDGDKESEVEDVET
						DSGNSPEDLRKEIMIGLQYQAEI
						PPYLGEYDGNEKDSPQPKKMT
						GVQNAKEVLST
<u> </u>		<u> </u>	l	<u> </u>		0.41444

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4368	34736	Α	4409	1	4485	
4369	34737	A	4410	2	927	IDHMIGHKASLNKFKKIEIISSTL SGHNGIKLEINSKRDLQNHANT RK\LNNLLLNEHWVKNEIKMEI LKFFELNDHNDTTYQNLWDTA KATF\LLRGKFTALNAYIKKTER AQTDILRSHVKELEKQEQTKPK PSRRKEITKIREELNEMETNKK KIQKINETKSRFFEQINKIDRSLA RLAKKRREKIQITSIRNKTGDTT TDTTEIQKIIQGYYEHLYAHKLE NLEEMDKFLEKYNPPSLNQEEL DTLNRTITSNKIEMVIKKLPTKK KSPGPNGFTAEFYQTFK\EELVP ILSILVHKTEKEGTLP
4370	34738	Α	4411	405	517	
4371	34740	A	4413	1	1197	MEISELNAKLRSQEKEKQNEIIK LQLEKLQHFQEEKNKEIAILRN TIRDLEQRLSVGKDSHLKRENE QLKISADLIKEKLKSHEQEYKN NIAKLVSEMKIKEEGYKKEISK LYQDMQRKGRIKVTCEWTCSE RKTEGREPGPVREPTGRSQSAE NEGSKTLAEINTKGTQSPAERIN KIDRLLARLTNKRREKVQISSIR NKTGDIRTDTTEKQKFMQGYH EHLYMHKLENLKEMDKFLEIY SPPRLKREDIETLSRPITISDIEM KNLKIPPKLPELINKFSKVSRYK INVHKLVALLYANSDQTDNQIK NSTHFTIVAKKYLGIYLTKDM KDLHKENSK/PLLKEIIDDTIKW KHIPCSWMSTTNIVKMTILPKTI YKFNAIIIKIPPSFFAERKKQS MIQRKRASIGAPCAWVRKKEE
4372	34740	A	4413	1	190	EEEEEEEEEEEEEEEEEK KKKKKKKERTTWLWGNPLT
4373	34741	A	4414	303	429	
4374	34742	Α	4415	123	252	
4375	34743	Α	4416	1	156	
4376	34744	A	4417	3	351	EEEEEEEEEEEEEEEEEEEEEEEEEEEEEEEEEEEEEE
4377	34745	Α	4418	1	192	
4378	34746	A	4419	3	259	
4379	34747	Α	4420	1	279	
4380	34748	В	4421	1	708	
4381	34749	A	4422	3	269	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4382	34750	A	4423	1	322	MAGKQGRSEGASSWRLSSVLQ LNSQYFLQGAQQCTFLAATAW KKRKEEEEEEEEEEEEEEEE EEEEEEEEEEEE'KKKKKKK KKKKKKKK
4383	34751	В	4424	327	674	
4384	34752	A	4425	494	960	TRFYDHALHLIHRKGSTTVRSP PPLYFIGESKASALLAISLRWSG RSQPRSSVNQIRKAWGFRPRKG TEE/DERSGCPSDALESDDPMA YIHFTAEGEVTFKSILFVPTSAP RGLFDEYGSKKSDYIKLYVRRV FITDDFHDMMPKYLNFVKGVV
4385	34753	A	4426		2539	VGGPRGWRCEDPNPGVGGGGG SCDRRGLETHRPHAMRALWVL GLCCVLLTFGSVRADDEVDVD GTVEEDLGKSREGSRTDDEVV QREEEAIQLDGLNASQIRELRE KSEKFAFQAEVNRMMKLIINSL YKNKEIFLRELISNASDALDKIR LISLTDENALSGNEELTVKIKCD KEKNLLHVTDTGVGMTREELV KNLGTIAKSGTSEFLNKMTEAQ EDGQSTSELIGQFGVGFYSAFL VADKVIVTSKHNNDTQHIWES DSNEFSVIADPRGNTLGRGTTIT LVLKEEASDYLELDTIKNLVKK YSQFINFPIYVWSSKTETVEEPM EEEAAKEEKEESDDEAAVEEE EEEKKPKTKKVEKTVWDWEL MNDIKPIWQRPSKEVEEDEYKA FYKSFSKESDDPMAYIHFTAEG EVTFK\SILFVPTSAPRG\LFD\DY GSKK\SDYIKLYV\RR\VFITD\DF HDMMPKYLNFVKGVVDSDDL PLNVSRETLQQHKLLKVIRKKL VP*NRWDMIKKI/SLDDKYNDT FW\KEFGYQHSSLVVIEGPLRIR TRLAKLLR\FQSSHHPTD\ITSLD QYVERMKEKQDKIYFMAGSSK KEAESSPFVERLLKKGYEVIYL TEPVDEYCIQALPEFDGKRFQN VAKEGVKFDESEKTKESREAVE KEFEPLLNWMKDKALKDKIEK AVVSQRLTESPCALVASQYGW

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4386	34754	Λ	4427	2	622	PARAALGILTSHQSGFLKTSTSK IT\STAWKN\KDITMQSTKQYAC LHDLTNKGIGEEIDNEHPWTKP VSSENFT\SP\YVWMLDAEDLA DIEDTVEWRHRNVESLCVMET ASNFSCS\TSGCFSKDIVG\LRTS\ ACWQQHCASPAFAYCG\HSFCC TGTALRTMSSLPESSAMW*KKP ARTRLPRGKDLIYFGSEKSDQE TGTLLLPVSS
4387	34755	A	4428	2	1421	QHCSQKDTAELLRGLSLWNHA EERQKFFKYSVDEKSDKEAEVS EHSTGITHLPPEVMLSIFSYLNP QELCRCSQVSMKWSQLTKTGS LWKHLYPVHWARGDWYSGPA TEL\DT\EPDDEWVKNR\KDESR AFHEWDEDADIDESEESAEESI AISIAQMEKRLLHGLIH\NVLPY VGTSVKTLVLAYSSAVSSKMV RQILELCPNLEHLDLTQTDISDS AFDSWSWLGCCQSLRHLDLSG CEKITDVALEKISRALGNSGHL HQSGFLKTSTSKITSTAWKNKD ITMQSTKQYACLHDLTNKGIGE EIDNEHPWTKPVSSENFTSPYV WMLDAEDLADIEDTVEWRHR NVESLCVMETASNFSCSTSGCF NHRPWSQNEYEQLNYAKQLKE RLEAFTRDFLPHMKEEEEVFQP MLMEYFTYEELKDIKKKVIAQ HCSQKDTAELLRGLSLWNHAE
4388	34756	В	4429	70	348	ERQKFFKYSVDEKSDKEAEVS
4389	34757	Α	4430	2	371	
4390	34758	A	4431		907	MGHRINIVCKIDAPCARQTRTF HPVVKTVEDCGRYPSVIEFGKY EIQTWYSSPYPQEYARNLAKEG KMGEREMSFVQQLQPMSGRCS LF\RELSSCTYLLNTQPP/AVSIH FLAVWIILLVDGNMSKIYCQNL CLLAKLFLDHKTLYYDVEPFLF YVLTKNDEKGCHLVGYFSKWT VLQGQWQVQGIAHFSRALTYLI CFSFPQEKLCQQKYNVSCIMIM PQHQRQGFGRFLIDFISFPRLTIG ASFTQLRKQSMSNST\EIPLLGD NGKSSPTFHWQSLTSSPNAHFS LEAQLSILGHLFQSP

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN 09/540,217	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	detection, (=possible nucleotide insertion)
4391	34759	A	4432	1	3468	MGKKQNRKTGNSKNQSASPPP
						KERSSSPATEQSWMENDFDELR
						EEGFRLSNYSELPEDIQTKGKE
						VENFEKNLEECITRITNKRNFKP
						TKIKRDKEGHYIMVKGSIQQEE
						LTILNIYAPTGAPRFIKQVLSDR
						QRDLDFHTLIMGDFNTPLSTLD
						RSTRQKVNKDTQELNSALHQA
						DLIDIYRTLHSKSTEYRFFSAPH
						HTYSKIDHLLGSKAFLSKCKRT
	}					EIITNYLSDHSAIKLELRIKNLTQ
						NRSTTWKLNN
4392	34760	Α	4433	3	1900	FNKCMTLKFRLKNFSRINKIDTP
		:				LARLIKKKREKNRIDTIKNDKG
						DITSNPTEIQSTIREYYKHLYTN
						KLENLEEMDKFLDTYTLPRLNO
						EEVESLNRPITGSEIMAIINSLPT
						KKSPGPDGFTAKFYQRYKEELV
						PFLLKLFQSIEKEGILPNSFYEAS
						IILIPKPGRDTTKNENFRPISLMN
	ŀ					IDAKILNKILANRIQQHIKKLIH
						HDQVGFIPGMQGWFNIRKSINV
						IQHISRTKDKNHMIISIDAEKAF
						DKIQQPFMLKTLNKLG\IKYLGI
						QLTRDVKDLFKERS/YEPLLNEI
						KEDTNKWKNIPCSWVGRINIVK
	ļ					MAILPKVIYRFNAIPIKLPMTFF
						TELEKTTLKFIWNQKRALIAKSI
		ŀ				LSQKNKAGGITLPDFKLYYKAT
						VTKTAWYWYQNRDIDQWNRT
			ĺ			EPSEITLHIYNYLIFDKPEKNKQ
						WGKDSLFNKWCWENWLAICR
				i		KLKLDPFLTPYTKINSRWIKDL
						NVRPKTIKTLEENLGITIQDIGM
						GKDYMSKTPKAMATKAKIDK
						WDLIKLKSFCTAKETTIRVNRQ
						PTKWEKIFATYSSDKGLISRIYN
						ELKQIYKKKTNNPIKKWVKDM
						NRHFSKEDIYAAKKHMKKCSP
						SLAIREMQIKTTMRYHLTPVRM
	1					AIIKKSGNN
4393	34761	A	4434	2	1932	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4394	34762	Α	4435	I	2571	MKAEIKMFFETNENKDTTYQN
						LWNTFKAMCRGKFIALNAHKR
		ľ				KQERSNTDTLTSQLKELKKQEQ
						THSKPSRRQEITKIRAEMKEIET
		ļ				QKTLQKIKESRTWFFEKINKIDR
		ŀ				LLARLTKKKREKNQIDAIKNDK
		İ				GDITTDPTEIQTTIREYYKHLYA
		İ				NKLENLEEMDKFLDTYTLPRLN
		ŀ				QEEVESLNRPITGSEIEAIINSLP
		ŀ				T/KKCPGPDGFTAEFYRRKRG\I
						LPNSFYEASIILIPKPGTDTTKKE
		İ				NFRPISLMNIDVKILNKILANRI
						QQHIKKLIHHDQVGFIPGMQG
		ŀ				WFNIRKSINIIQHINRAKDKNH
-					-	MIISIDAEKAFDKIQQCFMLKTL
İ		Ī				NKLGIDGTYLKIIRAIYDKPTAN
;		ŀ				IILNGQKLEVFPLKTGTRQGCPL
						SPLLFNIVLEVLARAIRQEKEIK
		ŀ				GIQLGNEEVKLSLFADDMIVYL
		ŀ				ENPIISAPNLLKLINNFSKGSAY
						KIKVQKSQAFLYTNNRQTESQI
		ļ				MSELPFTIASKRIKYLGIQLTRD
		ŀ	ŀ			VKDLFKENYKPLLKEIKEDTNK
						WKNIPCSWVGRINIMKMAILPK
		ŀ				VIYRFNAILIKLPMTFFTELEKST
ŀ						LKFIWNQKRARIAKSILSQKNK
						AGGITLPDFKLYYKATVTKTA
						WYWYQNRDIDQWNGTEPSEIM
						PHIYNYLIFDKPEKNKQWGKDS
						LFNKWCWENWLAICRKLKLDP
						FLTPYTKINSRWIKDLHVRPKTI
						KTLENLGNTIQDIGMGKDFMSK

SEQ ID	SEQ ID NO:		SEQ ID NO:		1	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		05/15/10,217	sequence	or pepade sequence	deterion, v possible nucleoniae moet non)
4395	34763	A	4436	1	1965	MTLESEQTFVYAVTATQTGAK
1333	31703	ļ'`	130	ľ	1703	EGTRMSKSNVAGQQGDSGEKA
						LQKTYQKILREKESALEAKYQA
}						MERAATFEHDRDKVKRQFKIF
		ŀ				RETKENEIQDLLRAKRELESKL
						QRLQAQGIQVFDPGESDSDDNC
						TDVTAAGTQCEYWTGGALGSE
						PSIGSMIQLQQSFRGPEFAHSSID
						VEGPFANVNRDDWDIAVASLL
l						QVTPLFSHSLWSNTVRCYLIYT
						DETQPEMDLFLKDYSPKLKRM
						CETMGYFFHAVYFPIDVENQYL
						TVRKWEIEKSSLVILFIHLTLPRI
						KYLGIQLTRDVKDLFKENYKPL
						LNEIKEDTNKWKNILCSWTGR
						NNVMKMATLPKVIYRFNAIPIK
						LPMTFFTELEKTTLKFIWNQKR
						AHIAKTILSEKNKAGGIMLPDF
1						KLYFKATVTKAAWYWCQNRD
		1		İ		IDQWNRTEASEITPHIYNHLIFD
						KPDKNKKWGKDSLFNKWCWE
						NWLAICRKLKLDPFLTPYTKIN
						SRWIKDLNVRPKTIKTLEENLG
ł.						NAIQDIGMGKDFMTKTPKAMA
					}	TKAKIDKWDLIKLKSFCMAKET
						PIGVNRQLTEWEKIFAIYPSDKG
				1		LISRIYKELKQTYKKKTNNPIEK
						LAKEMNRHLSKEDIYAANRHK
						KKCSSSLVIREMQIKTT/MRYHL
						TPVRMAIIKKSGNNRCWRGCG
4396	34764	Α	4437	300	476	PDLSLWLPLTFFPSFQLW*I*QL
						CVLELLFSRSIFVAFSVFPEFES
						WPALLGWGSSPG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
•	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			-	sequence		
4397	34765	Ā	4438	413	1689	QKLYKPERIKYLGIQLTRDVKD
						LFKENY/KLNEIKEDTNKRKNIP
						CSWVGRINILKMAILQKVIYRF
						NAIPIELPITFFTKLEKTTLRFIW
						NKKRVHIAKSIPSKKNKAGGIM
						LPDFKLYYKATITKTAWYLYQ
		ŀ				NRDIDQWNRTEALGITPHIYNH
						LIFDKPDKNKQRGKDSLFNKW
						CWENWLVICRKLKLDAFLTPY
						TKINSRWIKDLNIRPKTIKTLEE
						NLGNTIQGIGMGKDFMTKTPK
						AMATKAKIDKRDLIKLKSFCTA
						KETNIRVNRQPIEWEKIFAIYRS
						DKGLISRIYKELKQIYKKKTNN
						SIKKWAKDMNRHFSKEDIYAA
		1				NRHEKKWSPSLVTREMQIKTIM
						RYHLTPVRIMTIKMSGNNRCW
						RGYGEIGMLLHCWWECKLVQ
						ALWKTVWRFLKDLELEIPFDPV
1200	24766	<u> </u>	4420		2404	IPLLGIYPKDYT
4398 4399	34766 34767	A A	4439 4440	3	2404 1572	MLVSFVSLGSLCLQPGSQTLLE
4377	34707	^	14440		1372	KNRTVKPHVSFTLLPALSHVSE
						KNEAESMNSLIPPPPNLHTPAQ
						APFPLPTKEQDRSSSPATEQSW
		į				TENDFDELTEVGFRRSVITNSSK
						LKEDVRTHCKEAKNLEKRLHE
						WLTRINSVEKTLNDLKLKSMA
	ł					RELHDTCTSFNSRFDQVEERVS
						AIEDQTNEINNGENGTKLENTL
						QDIIQENFPNLARQANIQIQEIRR
						TPQRYSSRKATPRHIIVRFTKVE
						MKEKVLRAAREKVLEVLARAI
						SQEKEIKCTQLGKEEVKLSLFA
		1				DDMIVCLENPVVSDHNVLKLIS
ĺ	ŀ	ĺ				NFSKVSVYKINVQKSHAFLYTN
						NRQTESQIMSELPFTITTKRIKY
	-					LGIQLTRAVKDFFKEKYKPLLN
						EIKKDTNKWKNIPCSCIGRINIM
						KMAIVPKVIYGFNAIPIKLPRTF
						FTELEKTTLKFIWKKKGAKTILS
						IKNKAGGIMLPDFKLYYKATVT
]				KIAWYWYQNRYINQRNRTETS
						EITSHIYNHL/IFDKPDKNKKWG
						KDSLFNKWCWENWL
4400	34768	В	4441	1	1558	

sequ	uence	hod A	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
		A	09/540,217	1 ' '	of peptide sequence	deletion, \=possible nucleotide insertion\
4401 347	769	A			1	,
4401 347	769	Ā	t	sequence		
			4442	837	4329	TWKGTTDRSTRQKVNKDTQEL
						NSALHQADLIDIYRTLHPKSTE
) I			•			YTFF/LAPHHTYSKIDHIVGSKA
						LLSKCKRTEIITNYLSDHSAIKL
	ł					ELRIKNFTQSRSTTWKLNNLLL
	ł					NDYWVHNEMNAEIKMFFETNE
						NKDTTYQNLWDAFKAVCRGK
						FIALNAHKRKQERSKIDTLTSQL
						KELEKQEQTHSKASRRQEITKIR
						AELKEIETQKTLQKINESRSWFF
						ERITKSDRPLARLIKKKREKNQI
						DTIKNDKGDIT
4402 347	770	A	4443	1	816	MRRDYPVKAFTSRKREQHVQK
				ļ		VPSKKSRQVQRTERRFLETTPD
						LLYQKEKDLLLISSSKKQPRPGI
						ERHYMMTQGSIHQEDVAILK/V
						YTSNKRASKYIQQ/TLLEIKGKI/
						AHPQIVGDFNTPTSTIDRTIRQQI
						SIEFYDTIKQWDLTDTCRTGHPI
						TEYIFCSGAHLTFTKINHIQGPK
						RILKRFKRIEHECVLVLKGCQA
						KNRKKEEDLQTYWMLNIYGPH
						YRSGSYAAIHRQETICSGQLSQ
						ALRDRFAMNAKLLLSLAAHLW
						VIKLDFM
4403 347	77 l	A	4444	87	307	
4404 347	772	A	4445	1	534	MEESRGAKPPPALLPGDATLPP
						GSLGSARHPPEP/RPVPGP/PPHQ
						TCPGPSACSSRRPEPRSSPGSPA
	1					RAPPAPPPPAAPAPRCEPPLWLL
	l					LRVPCPGRSGWSWMTT*I/SERP
	ŀ					VQKRARSGPQPRLPPCLLPLSPP
						TAPDRATAVAT\PPVLGPMSSW
						SPRRAGGPTRPCTALQALSIPA
4405 347	773	A	4446	164	660	YPSGRRLREPADVADAWDGME
						ESRGAKPPPALLPGDATLPP\AP
						SGQLGTRPSPPSSRPSPHQTCPG
						PSACSSRRPEPRSSPGSPARAPP
						APPPPAAPAPRA/SPRRPLPAPRS
						ASVPAFSAPPSQWPEVGPSPCA
	İ					LRRAMPRGPGPPPEPRLVAEPG
						EDAAPTAGR
4406 347	774	A	4447	1	417	

SEQ ID NO: of peptide sequence			Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
34775	A	4448	1	1802	MSYPADDYESEAAYDPYAYPS DYDMHTGDPKQDLAYERQYE QQTYQVIPEVIKNFIQYFHKTVS
					DLIDQKVYELQASRVSSDVIDQ KVYEIQDIYENSWTKLTERFFK
					NTPWPEAEAIAPQVGNDAVFLI LYKELYYRHIYAKVSGGPSLEQ RFESYYNYCNLFNYILNADGPA
					PLELPNQWLWDIIDEFIYQFQSF SQYRCKTAKKSEEEIDFLRSNP
:					KIWNVHSVLNVLHSLVDKSNIN RQLEVYTSGGDPESVAGEYGR
					HSLYKMLGYFSLVGLLRLHSLL GDYYQAIKVLENIELNKKSMYS RVPECQVTTYYYVGFAYLMMR
					RYQDAİRVFANILLYIQRTKSM FQRTTYKYEMINKQNEQMHAL
					LAIALTMYPMRIDESIHLQLREK YG\DKMLRMQKGDPQVYEELF SYSCPKFL\SPVVPNYDNVHPN
					YHKE\PFLQ\QLKGVF**SSSQQ AQLS/TPIRSFLKLYT\TMP\VAK
					LAGFPGPSQSQEF\RIPGFFVFKQ QDERTSVWTQRVFSAPGW*NF
					SQASEVDF\YI\DKDMI\HIADTK VA\RRYG\DFFIRQI\HKF\EELNR TLKEGWGQRPWMIFHTHFREP GFECIIGQGSVFC
	of peptide	of peptide hod sequence	of peptide sequence hod in USSN 09/540,217	sequence 09/540,217 codon for peptide sequence	of peptide sequence hod in USSN location of first codon for last amino acid of peptide sequence of peptide sequence of peptide sequence

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IHHDHVSFJPRMQGWFNIHKP VIHHINRTINDKNHMISIDAME	SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
34776 A 4449 I 1722 MNIKAKILNKILANRIQOHIKK IHHDHYSFIPRMGGWFNIHKP VIHHINRTINDKNHMIISIDAEK FPKIQHPFTLKTLNKLDDMTV LENPIVSAQNILKLISNTSKYS YKINVQKSQAFLYTNNRQTES IMSELPTIJASKRIKYLGIQLTR VKOLFKENYKPLLKEIKEDTN WKNIPCSWVGRINVVKMAILP KVYYKFNAIPIKLPMTFFTELEK TTLKFIWNQKRARIAKSILSQN NKAGGITLPDFKLYYKATVTK AWYWYQNRDIDQWNRTEJPSI MPHINNHLTFDKPDKNKQWG DSLFNKWCWENWLAICRIKK DPFLTPYTKINSRWIKDLNVRR KTIKTLEENLGNTVQDIGMCK FMTKTPKAMATKAKIDKNDL KLSFCTSKETIIRVNRQPTEW KMFAIYPSDEGLISRICKEJPK YKKKNHPIKKWAKOMNRHF KEDIYVANKHMKKSLSSLVIR MQIKTTMRIHILTPVRMTIIKK GNRFWRGCGETGMLLAUD ECKLVQPL*KIVW*FLKDLESE PSDSAIPLGGIHPKAYKSFYY GWHGSPRCWR*PLPQRC GHLSCRWRTP GHKLLNKPGLKYKPYTNQNS GTMQNRAGFPRDEDCLLLQVI CHPYLTQEKLIQYCHSKGITYT AYSPLGSPDRPWARPEDPSLLI DPKIKEIAAKHKKTAAQVLIRF HIQRNVIVIPKSYTPARIVENIQ TEHYKYCGLCVGPNLEKNLY VVRMWKNSCGOPUL*ISSHLE DPFNAEY VVRMWKNSCGOPUL*ISSHLE DPFNAEY VVRMWKNSCGOPUL*ISSHLE DYPFNAEY VVRMWCMSCGOPUL*ISSHLE DYPFNAEY VVRMWCMSCGOPUL*ISSHLE DYPFNAEY VVRMWCMSCGOPUL*ISSHLE DYPFNAEY VVRMWCMSCGOPUL*ISSHLE DYPFNAEY VVRMWCMSCGOPUL*ISSHLE DYPFNAEY VVRMWCMSCGOPUL*ISSHLE DYPFNAEY DYPFNAEY DYPFNAEY DYPFNAEY DYPFNAEY DYPFNAEY DYPFNAEY DYPFNAEY DYPFNAEY DYPFNAEY	NO:	of peptide	hod		*		
4408 34776 A 4449 I 1722 MNIKAKILNKILANRIQQHIKK IHHDHYSFIPRMQGWFNIHKP VIHHINRTNDKNHMIISIDAEK FEKIQHPFTLKTLNKLDDMTV LENPVSAQNLLKLISNFSKVS YKINVQKSQAFLYTNNRQTES IMSELPFTIASKRIKYLGIQLTR VYOLFKENYKPLLEKIEDTN WKNIPCSWVGRINVVKMAILF KVIYRFNAIPIKLPMTFFTELEE TTLKFIWNQKRARIAKSNLSQN NKAGGITLPDFKLYYKATVTK AWYWYQNRDIDQWNRTEPS MPHIYNHLTFDKPDKNKQWG DSLFNKWCWENWLAICRKLK DPFLTPYTKINSRWIKDLNVRR KTIKTLEENLGNTVQDIGMCK FMTKTPKAMATKAKIDKWDL KLKSFCTSKETIIRVNRQPTEW KMFAIYPSDEGLISRICKE/LFK TYKKKNHPIKKWAKDNRHF! KEDIYVANKHMKSSLSSLVIR MQIKTTMRHHLTPVRMTIIKK GNNRFWRGCGETGMLHCW ECKLVQPL*RLVW*FLKDLESE PSDSAIPLGGIHPKAYKSFYY A4450 1050 1147 PGEWHOGGSPRCWFPLPQR GHLLSCRWETP A4410 34778 A 4451 I 614 MEELVDEGLVALGVSNFSHF QIEKLLNKPGLKVALGVSNFSHF QIEKLLNKPGLKVALGVSNFSHF QIEKLLNKPGLKVALGVSNFSHF QIEKLLNKPGLKVALGVSNFSHF QIEKLLNKPGLKVALGVSNFSHF QIEKLLNKPGLKVALGVSNFSHF QIEKLLNKPGLKVALGVSNFSHF QIEKLLNKPGLKVALGVSNFSHF QIEKLLNKPGLKVALGVSNFSHF QIEKLLNKPGLKKNLTPARIVENIQ TEHYKYCGLCVGPNLEKNLTF VDRM/WKNSCGGFVL*ISSHLE DPFNAEY VDRM/WKNSCGGFVL*ISSHLE DVFNAEY VDRM/WKNSCGGFVL*ISSHLE DVFNAEY VDRM/WKNSCGFVL*ISSHLE DVFNAEY VDRM/WKNSCGFVL*ISSHLE DVFNAEY VDRM/WKNSCGFVL*ISSHLE DVFNAEY VDRM/WKNSCGFVL*ISSHLE DVFNAEY VDRM/DDSSMHVSAPEDPPVG DVEAEDSDTDDPDPV		sequence		09/540,217		of peptide sequence	deletion, \=possible nucleotide insertion)
IHHDHVSFIPRMQGWFNIHKP VIHHINRTINDKNHMIISIDAM IHHINRTINDKNHMIISIDAM IHHINRTINDKNHMIISIDAM LENPIVSAQNLLKLISNFSKVS YKINVQKSQAFLYTNINQTIST IMSELPFTLASKRIKYLGIQLTR VKDLFKENYKPLLKEIKEDTN WKNIPCSWVGRINVVKMALL KVYRFNAIPIKLPMTFFTELER TTLKFIWNQKRARIAKSNLSQI NKAGGITLPPKLYYKATVIK AWYWYQNRDIDQWNRTEIPS MPHIYNHLTFDKPDKNKQWG DSLENKWCWENWLAICRILK DPFLITYTKINSRWIKDLINVR KTIKTLEENLGNTVQDIGMCK FMTKTPKAMATKAKIDKWDL KLKSFCTSKETIIRVNRQPTE KMFAJYSDGGLISSICKEFLEK KIYKKNHPIKKWAKDMNRHF KEDIYVANKHMKKSLSSLVIR KMGHTYBFDGELISSICKEFLEK KIYKKNHPIKKWAKDMNRHF KEDIYVANKHMKKSLSSLVIR MQIKTTMRHHLTPKNTHIKK GNNRFWRGCGETGMLLHCW ECKLVQPL*KIVW*FLKDLESE PSDSAIPLGGIHPKAYKSFYY 4409 34777 A 4450 1050 1147 PGEWHGQGSPKCWFLPQRC GHLLSCRWTP GHLLSCRWTP 4410 34778 A 4451 I 614 MEELVDEGLVKALGVSNFSHF QIEKLINKPGLKYKPYTNQNS GTMQNRAGPPRDEDCLLLQW CHPYLTQEKLIQYCHSKGITVT AYSPLGSPDRPWAKPEDPSLLE DPKIKEJAKHKKTAAQVLIR HIQRNVIVIPKSVTPARIVENILY VDRM/WKNSCGQFVL*ISSHLED DYPFNAEY 4411 34779 A 4452 2 240 WMELESLSHFQIEKLINKPGLKYKPVTNQNSIQFKGSILEEG VNMGDDSSMHVSAPEDPPVG DVEAEDSDTDDPDPV 4412 34780 A 4453 1 1019					sequence		·
IHHDHVSFIPRMQGWFNIHKP VIHHINRTINDKNHMIISIDAM IHHINRTINDKNHMIISIDAM IHHINRTINDKNHMIISIDAM LENPIVSAQNLLKLISNFSKVS YKINVQKSQAFLYTNINQTIST IMSELPFTLASKRIKYLGIQLTR VKDLFKENYKPLLKEIKEDTN WKNIPCSWVGRINVVKMALL KVYRFNAIPIKLPMTFFTELER TTLKFIWNQKRARIAKSNLSQI NKAGGITLPPKLYYKATVIK AWYWYQNRDIDQWNRTEIPS MPHIYNHLTFDKPDKNKQWG DSLENKWCWENWLAICRILK DPFLITYTKINSRWIKDLINVR KTIKTLEENLGNTVQDIGMCK FMTKTPKAMATKAKIDKWDL KLKSFCTSKETIIRVNRQPTE KMFAJYSDGGLISSICKEFLEK KIYKKNHPIKKWAKDMNRHF KEDIYVANKHMKKSLSSLVIR KMGHTYBFDGELISSICKEFLEK KIYKKNHPIKKWAKDMNRHF KEDIYVANKHMKKSLSSLVIR MQIKTTMRHHLTPKNTHIKK GNNRFWRGCGETGMLLHCW ECKLVQPL*KIVW*FLKDLESE PSDSAIPLGGIHPKAYKSFYY 4409 34777 A 4450 1050 1147 PGEWHGQGSPKCWFLPQRC GHLLSCRWTP GHLLSCRWTP 4410 34778 A 4451 I 614 MEELVDEGLVKALGVSNFSHF QIEKLINKPGLKYKPYTNQNS GTMQNRAGPPRDEDCLLLQW CHPYLTQEKLIQYCHSKGITVT AYSPLGSPDRPWAKPEDPSLLE DPKIKEJAKHKKTAAQVLIR HIQRNVIVIPKSVTPARIVENILY VDRM/WKNSCGQFVL*ISSHLED DYPFNAEY 4411 34779 A 4452 2 240 WMELESLSHFQIEKLINKPGLKYKPVTNQNSIQFKGSILEEG VNMGDDSSMHVSAPEDPPVG DVEAEDSDTDDPDPV 4412 34780 A 4453 1 1019	4408	34776	Α	4449	1	1722	MNIKAKILNKILANRIQQHIKKL
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PSDSAIPLGGIHPKAYKSFYY							
4409 34777 A 4450 1050 1147 PGEWHGQGSPRCWR*PLPQRO GHLLSCRWRTP 4410 34778 A 4451 I 614 MEELVDEGLVKALGVSNFSHF QIEKLLNKPGLKYKPVTNQNS GTMQNRAGFPRDEDCLLLQVE CHPYLTQEKLIQYCHSKGITVT AYSPLGSPDRPWAKPEDPSLLE DPKIKEIAAKHKKTAAQVLIRF HIQRNVIVIPKSVTPARIVENIQ TEHYKYCGLCVGPNLEKNLYF VDRM/WKNSCGQFVL*ISSHLE DYPFNAEY 4411 34779 A 4452 2 240 WMELESLSHFQIEKLLN/KPGL KYKPVTNQVNSIQFKGSILEEG VNMGDDSSMHVSAPEDPPVGG DVEAEDSDTDDPDV 4412 34780 A 4453 I 1019							-
GHLLSCRWRTP 4410 34778 A 4451 I 614 MEELVDEGLVKALGVSNFSHF QIEKLLNKPGLKYKPVTNQNS GTMQNRAGFPRDEDCLLLQVE CHPYLTQEKLIQYCHSKGITVT AYSPLGSPDRPWAKPEDPSLLE DPKIKEIAAKHKKTAAQVLIRF HIQRNVIVIPKSVTPARIVENIQ TEHYKYCGLCVGPNLEKNLYF VDRM/WKNSCGQFVL*ISSHLE DYPFNAEY 4411 34779 A 4452 2 240 WMELESLSHFQIEKLLN/KPGL KYKPVTNQVNSIQFKGSILEEG VNMGDDSSMHVSAPEDPPVGG DVEAEDSDTDDPDPV							
4410 34778 A 4451 I 614 MEELVDEGLVKALGVSNFSHF QIEKLLNKPGLKYKPVTNQNS GTMQNRAGFPRDEDCLLLQVF CHPYLTQEKLIQYCHSKGITVT AYSPLGSPDRPWAKPEDPSLLE DPKIKEIAAKHKKTAAQVLIRF HIQRNVIVIPKSVTPARIVENIQ TEHYKYCGLCVGPNLEKNLYF VDRM/WKNSCGQFVL*ISSHLE DYPFNAEY 4411 34779 A 4452 2 240 WMELESLSHFQIEKLLN/KPGL KYKPVTNQVNSIQFKGSILEEG VNMGDDSSMHVSAPEDPPVGG DVEAEDSDTDDPDPV 4412 34780 A 4453 I 1019	4409	34777	Α	4450	1050	1147	1
QIEKLLNKPGLKYKPVTNQNS. GTMQNRAGFPRDEDCLLLQVE CHPYLTQEKLIQYCHSKGITVT AYSPLGSPDRPWAKPEDPSLLE DPKIKEIAAKHKKTAAQVLIRF HIQRNVIVIPKSVTPARIVENIQ TEHYKYCGLCVGPNLEKNLYF VDRM/WKNSCGQFVL*ISSHLE DYPFNAEY 4411 34779 A 4452 2 240 WMELESLSHFQIEKLLN/KPGL KYKPVTNQVNSIQFKGSILEEG VNMGDDSSMHVSAPEDPPVG DVEAEDSDTDDPDPV 4412 34780 A 4453 1 1019							·
GTMQNRAGFPRDEDCLLLQVECHPYLTQEKLIQYCHSKGITVTAYSPLGSPDRPWAKPEDPSLLEDPKIKEIAAKHKKTAAQVLIRFHIQRNVIVIPKSVTPARIVENIQTEHYKYCGLCVGPNLEKNLYFVDRM/WKNSCGQFVL*ISSHLEDPFNAEY 4411 34779 A 4452 2 240 WMELESLSHFQIEKLLN/KPGLKYFVTNQVNSIQFKGSILEEGVNMGDDSSMHVSAPEDPPVGGDVEAEDSDTDDPDPV 4412 34780 A 4453 1 1019	4410	34778	Α	4451	1	614	ŀ
CHPYLTQEKLIQYCHSKGITVT AYSPLGSPDRPWAKPEDPSLLE DPKIKEIAAKHKKTAAQVLIRE HIQRNVIVIPKSVTPARIVENIQ TEHYKYCGLCVGPNLEKNLYE VDRM/WKNSCGQFVL*ISSHLE DYPFNAEY 4411 34779 A 4452 2 240 WMELESLSHFQIEKLLN/KPGL KYKPVTNQVNSIQFKGSILEEG VNMGDDSSMHVSAPEDPPVGG DVEAEDSDTDDPDPV 4412 34780 A 4453 1 1019	ļ	ļ					1 -
AYSPLGSPDRPWAKPEDPSLLE DPKIKEIAAKHKKTAAQVLIRE HIQRNVIVIPKSVTPARIVENIQ TEHYKYCGLCVGPNLEKNLYE VDRM/WKNSCGQFVL*ISSHLE DYPFNAEY 4411 34779 A 4452 2 240 WMELESLSHFQIEKLLN/KPGL KYKPVTNQVNSIQFKGSILEEG VNMGDDSSMHVSAPEDPPVG DVEAEDSDTDDPDPV 4412 34780 A 4453 1 1019							1
DPKIKEIAAKHKKTAAQVLIRF HIQRNVIVIPKSVTPARIVENIQ TEHYKYCGLCVGPNLEKNLYF VDRM/WKNSCGQFVL*ISSHLE DYPFNAEY 4411 34779 A 4452 2 240 WMELESLSHFQIEKLLN/KPGL KYKPVTNQVNSIQFKGSILEEG VNMGDDSSMHVSAPEDPPVGG DVEAEDSDTDDPDPV 4412 34780 A 4453 1 1019							CHPYLTQEKLIQYCHSKGITVT
HIQRNVIVIPKSVTPARIVENIQ TEHYKYCGLCVGPNLEKNLYF VDRM/WKNSCGQFVL*ISSHLE DYPFNAEY 4411 34779 A 4452 2 240 WMELESLSHFQIEKLLN/KPGL KYKPVTNQVNSIQFKGSILEEG VNMGDDSSMHVSAPEDPPVGG DVEAEDSDTDDPDPV 4412 34780 A 4453 1 1019							AYSPLGSPDRPWAKPEDPSLLE
TEHYKYCGLCVGPNLEKNLYF VDRM/WKNSCGQFVL*ISSHLE DYPFNAEY 4411 34779 A 4452 2 240 WMELESLSHFQIEKLLN/KPGL KYKPVTNQVNSIQFKGSILEEG VNMGDDSSMHVSAPEDPPVGG DVEAEDSDTDDPDPV 4412 34780 A 4453 1 1019							DPKIKEIAAKHKKTAAQVLIRF
VDRM/WKNSCGQFVL*ISSHLE DYPFNAEY 4411 34779 A 4452 2 240 WMELESLSHFQIEKLLN/KPGL KYKPVTNQVNSIQFKGSILEEG VNMGDDSSMHVSAPEDPPVGG DVEAEDSDTDDPDPV 4412 34780 A 4453 1 1019							HIQRNVIVIPKSVTPARIVENIQN
DYPFNAEY 4411 34779 A 4452 2 240 WMELESLSHFQIEKLLN/KPGL KYKPVTNQVNSIQFKGSILEEG VNMGDDSSMHVSAPEDPPVGG DVEAEDSDTDDPDPV 4412 34780 A 4453 1 1019							TEHYKYCGLCVGPNLEKNLYP
4411 34779 A 4452 2 240 WMELESLSHFQIEKLLN/KPGL KYKPVTNQVNSIQFKGSILEEG VNMGDDSSMHVSAPEDPPVGG DVEAEDSDTDDPDPV 4412 34780 A 4453 1 1019							VDRM/WKNSCGQFVL*ISSHLE
KYKPVTNQVNSIQFKGSILEEG VNMGDDSSMHVSAPEDPPVG DVEAEDSDTDDPDPV							DYPFNAEY
VNMGDDSSMHVSAPEDPPVG0 DVEAEDSDTDDPDPV 4412 34780 A 4453 1 1019	4411	34779	A	4452	2	240	WMELESLSHFQIEKLLN/KPGL
4412 34780 A 4453 1 1019 DVEAEDSDTDDPDPV							KYKPVTNQVNSIQFKGSILEEGI
4412 34780 A 4453 1 1019 DVEAEDSDTDDPDPV							VNMGDDSSMHVSAPEDPPVGQ
4412 34780 A 4453 1 1019							•
4413 34781 A 4454 1 2028	4412	34780	A	4453	1	1019	
[11] [2020] [1] [2020] [1]	4413	34781	A	4454	1	2028	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4414	34782	A	4455	3	1045	DFTSENFSAAWYLIENHSNTSF
1						EQLKMAVTNLKRQANKKSEGS
					ļ	LAYVKGGLSTFLEAQDALSAIH
						QKLEADGTEKVEGSMTQKLEN
				:		VLNRASNTADTLFQ/EKVLGRK
						DKADST\RNALNVL\QRFKFLFN
						LPLNIERNIQKGDYDVVINDYE
	ĺ					K\AKSLFG\KTEVQVF\KKYYAE
						V\EPRVEALRELLL\DKLLETPST
	1			l ,		LHDQKRYIRYLSDLHASGDPA
						WQCIGAQHKWILQLMHSCKEG
	i					YVKDLKGKDFSSNVFQFSGSAL
						RRVPDTVRVLDSQFSRSALRSV
						PDTVQVLDSQFSGSALRRVPDT
						VRVLDGQFSRSALRSVPDTVRV
				1		LDKCHCSPAKVVMNAVTIFTG
4415	34783	Ā	4456	1	440	MQRNLARAFSPGIKKIKMMCL
17-175	34703	` `	130		' ' '	GNSEKDWPKFRGVGEDAGLLA
				1		ARECGALLVIRHLINAVRAIVP
		l				NKSNNEIILVLQHFDNCVDK\TV
		1				QAFMEGSASEVLKEWTVTGKK
		1				KLLLQGEEELARLPFITGGSGSC
						YSSSTLAVEEECRVLA
4416	34784	A	4457	1	276	MEDEMEGLTEAGFRRWVTTNS
14410	34764	^	4437	'	270	AELKEHVLTQCKEAKNLDKRL
						EELLSRITSLERDISDQME/RELC
						EAYTSINSQINQAEERISEFEDH
						LAEI
4417	34785	A	4458	3	361	EMVHRKKKAVHRTATADDKK
7717	37703	^	1430		301	LQFSLKKLEVNNVSGIEEVNMF
		1				TNQGTVIHFN\AEMPANSFTITG
						HAETKQLMEMLPSILNQLGAH
						CLTSLRRLAEALPKQSVNGKAP
						LATGEDDDEVPA
4418	34786	Α	4459	1	475	BATTOEDDE VIA
4419	34787	A	4460	57	820	EDGSGGGKFPEGARQGGTGQR
'''		•		-	-20	RRRKAMRRTGAPAQADSRGRG
						RARGGCPGGEATLSQPPPRGGT
			1			RGQEPQMKETIMNQEKTRHTC
					:	RAQ\VRIGGKGTARRKKKVVH
						RGAAS/ADDKKLQF\SLKK\LGV
			}			NNISGIE\EVNM\FTN\QGTSGST
					Ì	FNNP*KFQGISWPANTFHHLQG
						HAEDKGS*QEMLAQHLKPSLG
						ADSLTSLRRLAEALPKQSVDGK
						APLATGEDDDDE\VPDLV\ENF*
						*RLPRNEANLNLSQLLKIKP
4420	24700	Α_	1161	1	1527	KLI KINEANLINLSQLEKIKP
4420	34788	Α	4461	1	1527	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
4421	34789	A	4462	8	327	LIWQLTFTKTIKS/CEEYGKIVST
						KAILDKNTNQCKGMCKGIRTL
						KSCLCYLINGSSIVEVQKRLAY
						AGTLEPSLVHQVYSELSYYKLP
						GTQVVRIIAEVLRMQDSSE
4422	34790	A	4463	2	573	WMEGREKWRGRRKDGRKEGR
						KEGRKERRREREKGRERK/GKE
						RKGKERKGKERKGRER
						KGTEGKGTEGKGKERKGKEGK
						GKERKGKERKGKEGKG
						KEQKGKERKGKERKGK
						ERKGKERKGKEEKGRERKGKE
						GKGRERKGKEGGKEEGRKERR
						KEGRKEGKKFSNNGMVEEMQ
4423	34791	В	4464	1	1344	
4424	34792	A	4465	3	373	
4425	34793	A	4466	1	3864	MQWEEAEKDPSGSCVFQRPPV
						ALVFPLHSKWTLVNSPPSSGDP
						YVPGRPAQSGQLSLSPAPPYVL
						PGPGKIKQAGNNPSLTSIYRSEV
						FCAHRHLHPPQLVCARGHIGSA
	ł					HLSVDRGSLIWEVLESTVWART
						NEWSPVTRTVLISALASTHIPQP
						CESRPPVPPEYEVTVLRSQGTA
						QLPPWSSSTSWRLTDPSCPKHA
						AWLTDLASSKGPAAGGTGSFS
						QPGTLTSTRTNPLKKEKSPEDL
						KQIKIDLGKFSDN
4426	34794	A	4467	3	415	RQIRIDLORI 3DIN
4427	34795	A	4468	396	676	LCFPYAERPDLQFLC*DLCARSP
1421	34173	^	1700	1370	070	YLLQAQKYLQEF*AIPHLDQQT
	ŀ					EPPDPSVSFYLLDCTLNCTAQH
				1		KTC*KKSIGL*EQNQQTLSSIPY
						SHT
4428	34796	A	4469	1	858	MEWEDNLPLELGRTVAKLLSD
14420	34/70	\\\^\	4407	1	030	HSQTPLGIQMFLLFSLSLRKSPL
				1	i	VCLSYLFNFRFTLESEVQHLSG
ŀ						1
	ľ					AITLTAWPKIPFLGIREAKSPRS
					<u> </u>	ENTRLATILEAGHRHLGTSVSK
						DHPVTFWRPRRDLQSDLKQIKI
						DLGKFSDNPDGYIDVLQELGQS
						FDLTWRDIMLLLNQTLTPNERS
						ATITAAREFGDLWYISQVVAAV
						AGLVSEAVKIIQG\LTVWT/SHD
				1		VNGILTAKGDLWLSDNHLLKY
						QALLLEGPVLRLRTCATLNPAT
						FLPDNEEKIEHNCQQVIAQTYA
4429	34797	Α	4470	918	1939	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4430	34798	A	4471	3	2693	PQVCLTIESQEVNCLLDAGAAF SVLLSCPGQLSSRSVTIRGVLGQ PVTRYFFQPLSCDWGALPFSHA FLIMPESLTPLLEREILVKAGAII HLNIGEGTPICRLLFEEGISPEV WATEGQYGQAKNAHFVQVKL KDSTSFPYQRQYPLRPEAQQRL QKIVKDLKAQGLVKPYSSPCNT
						PILGVQKPKRQWRLVQDLRIIN EAVFPLYPAIPSPYTLLSQIPEEA EWFTVLDLKDAFFCIPVHPDSQ FLFAFEDPSNPTSQLTWTVLPQ GFRDSPHLFGQALAQDLSQFSY LDTPVLQCMDDLLLAARSETLC
						HQATQALLNFLTTCGYKVSKP KAQLCSQQVKCLGLKLSKVTR ALSEERIQPILAYPYPKTLKQLR GFLGITGFCRIWIPRYGKIARPL YTLIKETQKANTHLVRWTPEAE AAFHALKKALMQAPVLSLLTG QDFSSYVTKNKQTKKKK\T*IA
						LRVLALV*GTSLQPVAYLSKKT DVVAKGWPHCLWVMAAIAVLI SKAVKMIQ*RDLTVWTSHDVN GILTAKGDLWLSDNHLLKYQA LLLEGPMLRLCTCAALNLDTFL
						PHNEEKIEHNCQQVIAQTYATR GDHLEVPLTDPNPNLYTDGRSF VEKGLQKVGYAVVSDNGILES NPLTPGTSAQLAELIALTWALE LGEGKRVNIYTDSKYAYLVLH AHAVIWREREFLTSEGTPIKHQ
4431	34799	С	4472	11	1639	
4432	34800	A	4473	95	2539	
4433	34801	A	4474	345	768	PRGARSTRCLPVERR\CDGLQD CGDGSDEAGCPDLACGRRLGSF YGSFASPDLFGAARGPSDIHCT WLVDTQDSRRVLLQLELRLGY DDYVQVYEGLGERGDRLLQTL SYRSNHRPVSLEAAQGRLTVA YHARARSHPLMNE
4434	34802	A	4475	47	563	RLRFAVFTGAFHALSFLLSFVV LCCTYLKGLKVARFHCKRIDV/ MHHADARAAGGPAPQCAGTLS \EEQKRRRQRATKKISTFIGTFL VCFAPYVITRLVELFSTVPIGSH WGVLSKCLAYSKAASDPFVYS LLRHQYRKSCKEILNRLLHRRSI HSSGLTGDSHSQNILPVSE

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4435	34803	A	4476	1	332	ERGRQEMSAKLRPPAEPPCVPA RISP*RPS*RQ*MERRCPPWRCS PMPC/CFFREHALQVRCGPTSA DCGRDPLFSPHPKPLPHPVPDIG WVATAGAQRSSSPVPSSLFVW
4436	34804	A	4477	297	943	TGSWGGGADQLRPALTTALM PPDNRFGENTPAAPANGHCAP\ EPDITLVQDHSELPIGAAATMA HEIGHSLGLSHDPDGCCVEAAA ESGGCVMAAAT/GVRGHPFPRV FS/SCSRRQLRAFFRKGGGACLS NAPD/TRTPGAAALCGNGFVEA GEECYCVS\GQECRDLCCFAHN CSLRPGAQCAHGDCCVRCLVR/ CMEGSGSHQLPRLVPGGDSAEI LM
4437	34805	Α	4478	1	836	MGPLTFRDVKIEFSLEEWQCLD TAPGNLYRDVMLENYRNLVFL VMCSHFAQDVWPEHSIKDSFQ KVILRTYGKYGHENLQLRKDH KSVDACKVYKGGYNGLNQCLT TTDSKIFQCDKYVKVFHKFPNV NRNKIRHTGKKPFKCKNRGKSF CMLSQLTQHKKIHTREYSYKCE ECGKAFNWSSTLTKHKIIHTGE KPYKCEECGKAFNRSSNLTKH KIIHTGEKPYKCEECG\KAFNRS STLTKHKRIHTEEKPYKCEECG KAFNQFSILNKHKRIHMGR
4438	34806	A	4479	1	588	MLGKVQQQEQTIAKDLVVTKY KMCGGT/DIANRVLRSLVEASS S\GGQDYILKEGDLVKIDLGVH VDGFIANVTHTFVVDVAQGTQ VTGRKGDVIKAAQLCVEAALC LVKPGNQNIQVREAWSKVALS FNCMPIEGMLSHQLKQHVIDGE KNIIQNPTDQQKKDHEKAEFEV HEVYAADVLVSSGEGKAKDAG
4439	34807	A	4480	85	561	LSHCLPLQTTEVGGFGNLLGY WIACSIGCVLSTGMLSHQLKQH VIDGEKTIIQNPTDQQKKDHEK AEFEVHEVYAVDVLVSSGEGK VRRVPELAKRGD*ECSPDQMLL KLLFQAKDAGQRTTIYKRDPSK QYGLKMKTSRAFFSEVERRFD AMPFTLRY

SEQ ID	SEQ ID NO:		SEQ ID NO:	Nucleotide location of first		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO:	of peptide sequence	hod	in USSN 09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			,	sequence		
	<u> </u>			<u></u>		
4440	34808	Α	4481	1	1408	GTSAPQPARSQLLALACLPAPL
						LARAFARPLLEDRGDSDHSLW
						LGRETEAAAAQGKRGCSGGSR
						KMSGEDEQQEQTIVD/DSLVVT
						KYKMGGDIANRVLRSLVEASSS
						GVSVLSLCEKGDAMIMEETGKI
						FKKEKEMKKGIAFPTSIS\VNNC
						VCHFSP/L*KSDQDYILKEGDLV
						KIDLGVH\VDGFIANVAHTFVV
						DVAQGTQVTGRK\ADVIKAAH
						L\CAEAA\LRLVKPGNQNTQVT\
						EAWNKVAHSF\NCTPI\EGMLSH
						SLKQHVIDGEKP*FQNPTDKQK\
						RAHEKADFEVH\DVYAVEGLV
						KPQERARPKDAGQRTTIYKRDP
						SKQYGLKMKTSRAFFSEVERRF
						DAMPFTLRAFEDEKKARMGV\
						VECAKHEL/VWQPFNVLYSGRE
1						GDFVCPVLNFTVL\LMPNGPML
						ITSGPFEPDLYKSQMEVQ\DAEL
						KALLQSSASRKTQKKKKKKAS
						KTAENATSGETLEENEAGD
4441	34809	Α	4482	3	190	
4442	34810	В	4483	1	588	
4443	34811	Α	4484	1	1312	MSSKGSVVLAYSGGLDTSCILV
						WLKEQGYDVIAYLANIGQKED
		l				FEEARKKALKLGAKKVFIEDVS
						REFVEEFIWPAIQSSALYEDRYL
					•	LGTFF\ARPCIARKQVEI\AQREG
						AKYVSHGATGKGNDQVRFELS
						CYSLAP\QIKVIAPWRMPEFYNR
						FKGRNDLMEYAK\QHGIPIPVTP
						KNPWSMDENLMHISYEAGILE
						NPKNQAPPGLYTKTQDPAKAP
						NTPDILEIEF\KKGVPVEGGPTF
						KDG\TTHQTFL\ELF\MYLNEVA
						GKHGVGPYLTSWENRFHWELK
						SRGILRRPQAG\TILYHAHLDIE
						AFTMGGDRAQIPNQGLGFEFVE
						LGVYRFSGTAPECELVGPCLRQ
						SPQERVEGKSAGVPSLKGPRCT
						SLGPEVPHCSLYNE\ELVKHGT
						CQGDYE\PN*LPPGFIQTSISLKA
						EGNYHRLPRAKVTAQIRPRVQ
4444	34812	В	4485	47	482	

SEQ ID NO:	of peptide sequence		SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4445	34813	A	4486	2328	3435	KTTTLEDNLGNTIQDIGPGKDF MMKIPKANATKIKIDEWDLIKL KSFCTAKATTKRVNKHDESLRS HYE*WGMLTDCVVMRDPNTK RSRGCGFVTYATVEEVDAATN ARPHKVDGKVVEPKRTVSRED SQRPGAHLTVKKIFVGGIKEDT GGFAFVTFDDHDSVDKIVIPKY HTVNGHNCEVRKALSKQEMAS ASSSQRGRSGRGGGFGGNENFG CGGNFSGHGGFGGSHDGGGYG GSGDGYNGFGNDGGYPGGPG YSGGSRGYGSGGQGCGNQDSG YGRSGSYDSCNKGGRGGFGSG SGSNFGGGGSYNDFGNYNNQY SNFGPMKGGNF/GGRRSGP*GD GGQYFAKPPNHSGYGGSSSSSSS
4446	34814	Α	4487	1	762	000 11 ART 1 M130 1 00333333
4447	34815	Α	4488	3	333	
4448	34816	A	4489		1676	MRDPNTKRSRGFGFVTYATVE EVDAAMNTTPHKVDGRVVEPK RAVSREDSQRPGAHLTVKKIFV GGIKEDTEEHHLRDYFEQYGKI EVIEIMTDRGSGKKRGFAFVTF DDHDSVDKIVIQKYHTVKGHN CEVRKALPKQEMASASSSQRG RRGSGNFGGGRGDGFGGNDNF GRGGNFSGRGGFGGSCGGGY GGSGDGYNGFGNDGSNF*G\GG SYNDSGNYNNQSSKFEPMKGG NFGGRSSGPYGGGQYFAKPQ NQAARCVAARWLFRTEARLVF LQKFPPWAVVEVVTVIVAAPA AATATTRDGGGCSRNCNIPEVF PELLGCPNRRGPPGGVREKQQQ TNSKSTRQEITKTIAELKEIETR KTLQKINESRSWFYEKINKVDR LLDRLIKKKREKSQIDAIKNDIG DIVTDPAEIQTTIKEYYKRLYAN ELENLEEMDKFLATYSLHSLNQ EEVESLNKPVTSSEVEAVTNSL PTKKSPGPDGFTVLLEVLAPAIR QEKEIKHIPIGREEVKLSLFADD VIVYLENPIVSAQNLLKLISNFS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4449	34817	A	4490	1	11445	 MKCLKFINHKEILEASERKQAE
4449	34617	l ^A	4490	[I	1443	SLDFPFKKLRWHLCEGWIEEER
				[DESRKSETIFKOLFKVPVLKETI
•						YYKFYGPPVYQIETVYFMALSP
]		
	l					PKSKQFDKTKQNNNNKKTHQF
						VIVFFKTDEHLSARGRRRRSIVK
						VSLLPAVIGLKSKFLKKPDQLR
						KLF\IGGLSFETTDESLRSHFEQ
				ĺ		WGTLTDCVVMRDPNTKRSRGF
						GFVTYAT\VEEVDAAMNARPH
						KVD\GRVV\EPKRAVSREDSQRP
						GAHI/TLVKKIFVGGIKEDTEEH
						HLRDDYFEEIILNSMEKIEV\IEI
						MT\DRGSGKKRGFAF\VTFDDH
	1					DSVDKIVI\QKYHT/VGNGHNCE
						V\RKALSKQEDG*VLHPAQRG\
		ļ				RSGSGKLLVVGRGRWFSVGMD
						NFG\RGGNFSWSVVAFGGT\RG\
						GGGYGWQWGMAYNGFGNDG\
	ĺ					SNFGGGG\SYNDFG\NYNNQ\SS
						NFGPMKGGNFG\GRSSGPYG\G
		<u>L</u>				GGQYF\AKPR\NQGGYGGSSSS\
4450	34818	Α	4491	134	612	TVLNSMSVILAISTLLKIITGELL
						QSFGDGLLWNLVIGIRGIDGLSP
						KVRKVLQLLRLRQIFN/GTFVK
						LIKVTVNMLRTVEPYIAWGYPN
						LKSVNELIYKHGYGKISKKRIA
	1	ŀ				LTDNVLIARSLGKYGIICMEDLI
						YEIYTVGKRFKEANNFLWPFKL
4451	34819	A	4492	1	1983	
4452	34820	Α	4493	1	1527	
4453	34821	В	4494	1	2211	
4454	34822	Α	4495	1	2478	
4455	34823	Α	4496	2	1544	
4456	34824	В	4497	1	2151	
4457	34825	Α	4498	1	744	
4458	34826	В	4499	1	2172	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4459	34827	A	4500	1	2535	MKSGHPEKEQDNSDVQETREIT
		ŀ				IRGLLCTALMRHSTGAIAYLGV
						LSGSASLKLAGVPLRCCEGDKD
		ľ				AGHPLETQTALCERGRGARSLV
						GNTIMTSQPVPNETIIVLPSNVIN
				1		FSQAEKPEPTNQGQDSLKKHLH
				i		AEIKVIGVNLIQNVLERGWGKC
		1				QEMIYVLGLDICRPFFVSRVSEE
	İ	l				GRMGQRGEEDANSLDFPPASLL
						CLICQEQGVNGESCSPVGMYH
						REIVPVYEVLSVITGLQIQVFSG
						KEADSVIKRSIGWGPFFKPRTK
						DKNHMIISIDAEKAFDKIQQHF
						MLKTLSKLGIDGTYLKIIRAIYD
						KPTANIILNGQKLEAFPLKTGTR
						QGCPLSPLLFNIVLEVLARAIRQ
						EKEIKGIQLGKEEVKLSLFADD
	ļ			•		MIVYLENPIVSDQNLLKLISNFS
						KVSGYKINVQKSQAFLYTNNR
						QTESQIMSELPFTIASKRIKYLGI
						QLTRDVKDLFKENYKPLLNEIK
						EDTNKWKNIPCSWVGRINIVK
						MAILPKVIYRFNAIPIKLPMTFF
						TELEKTTLKFIWYQKRARITKSI
						LSQRNKAGDITLPDFKLYYKAT
						VNKTAWYWHQNRHIDQWNRT
						KPSEITLHIYNYLFFDNPDKNKK
				i		WGKDSLFNKWCWENWLAICR
		Ì				KLKLDPFLTPYTKINSRWIKDL
				1		NIRPKTIKTLEENLGITIQDIGMG
						KYFMTKTPKAMATKAKIDKW
						DLIKLKSFCTGKETTIRVNRQPT
4460	34828	В	4501	I	1785	
4461	34829	Α	4502	1	1415	
4462	34830	В	4503	1	3262	
4463	34831	Α	4504	1	278	

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deterion, (=possible nucleotide insertion)
4464	34832	Α	4505	[3	2528	ENKDTTYQNLWDAFKA\VCRG
						KFIALNAHKRKQEKSKIDTLTS
İ		ľ				QLKELEKQEQTHSKASRRQEIT
ł						KIRAELKEIDTQKTLQKINESRS
						WFFERINKIDRPLARLIKKKREK
						NQTDTIKNCKGDITTDPTEIQTT
1						IREYYKHLYANKLENLEEMDK
		į				FLNTYTLPRLNQEEVESLNRPIT
						GAEIVAIISSLPT/K/KSPGPDGFT
				ł		AEFYQRYKEE/LEKEGILPNSFY
						EASIILIPKPGRDATKKENFRPIS
:						LMNIDAKILNKILAKRIQQHIKK
						LIHHDQVGFIPGMQGWFNIRKS
						INVIQHINRTKDKNHMIISIDAE
	1					KAFDKIQQRFLLKTLNKLGIDG
1						TYFKIIRAIYDKPTANIILNGQKL
1	:					EAFPLKTGTRQGCPLSPLLFNIV
		İ				LEVLARAIRQEKEIKGIQLGKEE
		1				VKLSLFADDMIVYLENPIVSAQ
						NLLKLISNFSKVSGYKINVQKS
						QAFLYTNNTQTESQIMSELPFTI
		1				ASKRIKYLGIQLTRDVKDLFKE
		ł				NYKPLLKEIKDDTNKWKNIPCS
						WVGRINIVKMAILPKLPMTFFT
						ELEK\TTLKFIWNQKRACIAKSI
						LSQKNKAGGITLPDFKLYYKAT
		l				VTKTAWYWYQNRDIDQWNRT
		l				EPSEIMPPIYNYLIFDKPEKNKQ
						WGKDSLFNKWCWENWLAICR
		ŀ				KLKLDPFLTPYTKINSRWIKDL
		ŀ				NVRPKTIKTLEENLGITIQDIGL
						GKDFMSKTPKAMATKAKIDK
4465	34833	В	4506	1	5401	
4466	34834	Α	4507	1	5271	MNIDAKILNKILPNQIQQHIKKL
						IHHDQVGFIPGMQGWFNIRKSI
İ				:		NVIQHINRAKDKNHMIILIDAEK
				1		SFDKIQQPFMLKTLNKLGIDGT
						YFKIIRAIYDKPTANIILNGQKLE
						VFTLKTGTRQGCPLSPLLFNIVL
						EVLARAIRQEKEIKGIQLGKEEV
						KLSLFADDMIVYLENPIVSAQN
				1		LLKQISNFSKISGYKINVQKSQA
						FLYTNNRQTESQIMSEIPFTIAL
] .		KRIKYLGIQLTRDVKDLFKENY
4467	34835	В	4508	924	3423	
4468	34836	Α	4509	525	673	RDSWGTCPVSGAGKVDWPPSS
						*HHR*HQQWCCGMPHQLSTKE
						NISIKDHLTKEKRKGGAV*RII

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	j	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
4469	34837	A	4510	25	11766	GTCQFAAMNVVFAVKQYISKM
.,05	15.057					IEDSGPGMKVLLMDKETTGIVS
						MVYTQSEILQKEVYLFERIDSQ
		1				NREIMKHLKAICFLRPTKENVD
				•	İ	YIIQELRRPKYTIYFIYFSNVI\SK
		ļ		}		SDVEVIGLKLIEQEVVAEVQEF
		1				YGDYIAVNPHLFSLNILGCCQG
						•
		1				RNWDPAQLSRTTQGLTALLLSL
						KKCPMIRYQLSSEAAKRLAECV
						KQVITKEYELFEFRRTEVPPLLL
						ILDRLDDAITPLLNQWTYQAM
						VHELLGINNNRIDLSRVPGISKD
Í						LREVVLSAENDEFYANNMYLN
						FAEIGSNIKNLMEDFQKKKPKE
						QQKLESIGS\MKA\FVENYPQFK
						KMSGTVSKHVTVVGELSRLVS
						ERNLLEVSEVEQELACQNDHSS
						ALQNIKRLLQNPKVTEFDAARL
		1				VMLYALHYERHSSNSLPGLM
		-				MDLRNKGVSEKYRKLVSAVVE
	į					YGGKRVRGSDLFSPKDAVAITK
					İ	QFLKGLKGVGNV\YTQLQPF\L
						H\ETLDHLIKGRLKENLYPYLGP
						STLRDRPQDIIVFVIGGATYEEA
						LTVYNLNRTTPGVRIVLGGTTV
						HNTKSFLEEVLASGLHSRSKES
4470	34838	Α	4511	1	1335	MAPVTMMGYRSGKMGILADV
	ŀ					QLQVGPPGPWLHLVVIAPVPEC
						ITGIGIFSSWGSPDVGPLLYDIR
	ļ					AIMWGSLAPAENTWILGNNHR
	ŀ					RFLAQLKPRVIMQDFSNVISKS
	ļ	-				DVKSLAEADEQEVVAEVQQVI
						TKEYELFEFRRTEVPPLLLILDR
						CDDAITPLLNQWTYQAMVHEL
	ŀ			1		LGINNNRIDLSRVPGISKDLREV
	ļ					\VSSAEIDEFYANNMYLNFAEIG
	ŀ		1			SNIKNLMEDFQKKKPKEQQKL
						ESIADMKAFVENYPQFKKMSG\
	ļ					TVSK\HVTVVG\ELSRL\VSERN
						LAGRFSEVEARNWAC\QNDHS\
						SALQNIKRLLQNPKVTEFDAAR
		l				LVML\YA\LHYERHSSNSLP\GL\
						MMDLRN\KGVFWRKYSKARVL
						AVVEYGGKRVRGSDLFSPKDA
						VAITKQFLKGLKQQEIVNCVLA
						AANVYIKQLPLSIQPSASLNGCI
	:					SLEKKPLVSTQRN
4471	34839	A	4512	1	816	DEDIKIC DIDIKICI
4471	34840	A	4513	26	257	
4473	34841	A	4514	56	236	
77/3	124041	112	1-10 1-1	150	230	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence		in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4474	34842	Α	4515	170	373	HSPRGSTASF/CEVSETKNPPIPD TPATREAEGLRSRLAVYSTRDS PCVACSGSYTQAQGSLGRKFQ DP
4475	34843	A٠	4516	262	358	
4476	34844	A	4517	2298	2556	NHKNPRRKPRQYHSGHRHGQG LHD*NTKSNGTKSNGNKSQN* Q\WDLINLKSFCTAKETTIRVNR QPTEWEKIFTIYPSDKGLISRI
4477	34845	Α	4518	801	944	DQEPTNSRHILATQMGPSPITKQ SNPGV**KECGFSCSPRVWRYL VS
4478	34846	В	4519	85	660	
4479	34847	Α	4520	693	827	
4480	34848	Α	4521	272	339	
4481	34849	С	4522	532	2754	
4482	34850	В	4523	1	519	
4483	34851	В	4524	266	935	
4484	34852	Α	4525	1	1584	
4485	34854	A	4526		335	GALPNGDRGRRKSRFALYKRP KANGVKPSTVHVISTPQASKAI SCKGQHSISYTLSRNQTVVVEY THDKDTDMFQVGRSTESPIDFV VTDTISGSQNTDEAQITQSTISR FACRIVCDRNEPYTARIFAAGF DSSKNIFLGEKAAKWKNPDGH MDGLTTNGVLVMHPRGGFTEE SQPGVWREISVCGDVYTLRETR SAQQRGKLGLQTGDMAENT/T VHALPSNCMVWRRSQTRQQIS
4487	34855	A	4528	328	871	DCGGGRARTAIFAGAARAADN
440/	34633		J+320			KKCAGARRALGRARGCSATAR PRRRRRPRGLAPPRPARPPPG GMSYKPNLAAHMPAAALNAA GSVHSPSTSMATSSQYRQLLSD YGPPSLGYTQGTGNSQVPQSKY AELLAIIEELGKEIRPTYAGSKSS \MERL\KRGIIHARGLVRECLAE TERNARS
4488	34856	Α	4529		653	MAGPAESSPQGAHPNSPFALQH HSSLTVKPLHRQNVIQHQVAG QENRRGHQAGSSTSPQPLEALK RPNLRAPFHSQSRRILIPPAGNP TPGAAAPADPSTQRRDRWGCA LPMPRVAAGSAHHQQAGPTAA AQHRTPVALFSPPLSLVYGQQQ RKESETPTVPTPPARARGWTET GVEHVPAYNRTRAPEKCDI/SV PSPHSSPDAETSHPRHISPCPG

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4489	34857	A	4530	3	432	NSRVDDFVAAQDAKGKKVAP APAVVKKQEAKKVVNPLFEKR PKNFGIGQ\QRLLARAEKKAAG KGDVPTKRPPVLRAGVNTVTT LVENKKAQLVVIAHDVDPIELV VFLPALCRKMGVPYCIIKGKAR LGRLVHRKTCTTVAFT
4490	34858	A	4531		2073	MKPCAHSWNAELSRNIIRHSFN LVMVAASQVAVSQLLGSYEILL LVSIELMFCFGLGYFFIPMQEW PNTYGERVFVDVESSVFKWNH KCLHKTEAERDYTKKRLKLCG HKPGNAVGQQKLEEARNRFFT RAPGGSAALPTLRFQPSDTDFR LLASRTILTFETKNPSELAERLR SVCGNQSNAYARLLEYRLNAL RGLWNAQRQLALEEQHERESS GDEETLALLKRQGLLQQPEQAP FTSRMGLLLVFPLIQSQSRTDPS LCNITAEVLLNCLRDCQPLSLT KEPADCLNGIETLLCSWLEETS DTGRHIPHKQKENAAAALVAL ACARGFVYCRNEELEPGWVAF GSGSLLHRPVSFDNKPHSLFQVI DQNTLQVCQVVPMPANHLPIG STMSTVHLSSDGTYFYWIWSPA SLNEKTPKGHSVFMDIFELVTL KGKKAKGKKVAPAPAVVKKQ EAKKVVNSLFEKR\DIQPKRELT YFVKW/PRYVRLQQQRAILYKQ LKVPPAINQFTQALNCQTVTQL LKLAHKYRPETKQEKKQRLLA QAEKKAAGKGGVPTKRPPALR AGVNTITTLVENKKAQLVVIAH DVDSIELVVFLPALCCKMGVPY CIIKGKARLGRLVHRKTCTTVA FTQVNLEDKGALAKLVEGIRTN DNDRYDEICCHWGGNILGPKS
1401	10.050	<u> </u>	4500		0565	VACIAKLEKAKAKELATKLG
4491	34859	A	4532 4533		2565 644	MPKGKKAKEKKVAPAPAVVK KQEAKKVVNPLFEKRPKNFGT GQDIQPKRDLTHFVKWPCYIRL QQQRTILYKWLKVPPEINQFTQ APDSQTATLLLKLAH/KYRPET NQEKKQRLLARAKKKAAGKG DIP\TKSPPVLRAGVNTITTLVE NKKAQLVVIAHDVDPIKLVVFL PVLCHK/MGVPYCIIKGKARLG HLVHRKTCTTVTFTQVNSEDK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
4493	34861	Α	4534	1	931	KSIQKGLKMCLSSSLLPPSKMP
						KGKKAKGKKVAPAPAVVKKQ
						EAKKVVNPLFEKRP\KNFGIGQ
						DIQP\KRDFTRFVKWPRLLSGC
						MR\KRAILYKAG*KLPPA\INQF
						HPGPWDPANKLLQLL*AWAHK
1						\YRP\ETKAKRKKQRL\LARA\E
1			ľ			KKAA\GKGDVPNERDPPV\LRA\
		ŀ				GVNTVTHLWWRNKKAPAWVV
						IATRRWIPFEL\VVFLPAL\CREK
						WGSPYCIIKGKARLGR\LVH\RR
						PCT\TVGFHTR*NSKDKRRLLA*
	1					AGLEAIRTQFTIDQIRWRSGRH\
						WG\GNVLG\PKSVARIRQASKR
						QRLKELATKLG
4494	34862	A	4535	3	227	
4495	34863	Α	4536	1	338	
4496	34864	Α	4537	1	352	
4497	34865	Α	4538	2	368	
4498	34866	A	4539	3	468	DD CDNDDD VTCOCD DATE NICOD
4499	34867	Α	4540	2	790	PRGRNRRRKTFQERRMTLNESP
						EKIGKWIECYGHPPASKLVEIYI HTVFVEDKLSICIRSFNKKADGS
						WRMTVDYCKLNQVVTAIAAAI
						PDVVSLLEQINTSPDTWYAAID
}						LANALFSIPVHKGYINSLALCH
}						NVIWRELDCFSLPRDTTLVHYI
						DDIMLIGSSVQEVENKLDLLVK
						DKLLHLAPPTTKEEVQHMVGL
						FGFWRQHIPHLGVLHQPIYRVI
	ŀ					RKAA/SFEWGPEQEKALQQVQ
						AAVGGKQSENNLGHQRSPGLW
4500	34868	В	4541	179	1219	
4501	34869	Α	4542	1706	2517	THLLVPGMQPLTWQMPFSPFLS
				İ		ISPTRSNLPSAATPVIAQWA/HE
	ŀ					QSGHGGRDGGYTWAQQHGLA
		İ	}			FTNTDLATVNAKIGFAYPVCDA
						SAKTTIRGLLECLIRCDGIPHSIA
						SDQARIHRSRNQEVEVEVAPLT
						ITPSDPLAKFLLSVPVTLRSAGL
ŀ						EVLVPGEGMLPPGNTRTIPLNW
						KLRLPPGHFGLLLTLSQEAKNG
1						VTVLAGVIDLDYQDEISLLLHN
		Ì				GGKKEYARNTGDPLGRLLVLP
[CPVIKINGKLQQPNPGGTTNGS
						DPSGMKV
			<u> </u>			

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4502	34870	A	4543	3	367	DLWPFTRVTVH/WGKANDQTF
.502	13.070					QGLLDTGSELTLIPGYPKRHCC
						PPVKVRVYGGQTDGSWRMTV
						GYHKLNOVVTPIAAAVPDVVS
						LLEQINTTPAIKWVVHSSIPSSN
						GSGVYVIRLEQVLKAQ
4503	34871	Α	4544	2	541	
4504	34872	В	4545	1	681	
4505	34873	A	4546	2	1091	PRGRNRRRKTFQERRMTLNESP
						EKIGKWIECYGHPPASKLVEIYI
						HTVFVEDKLSICIRSFNKKADGS
		-				WRMTVDYCKLNQVVTAIAAAI
						PDVVSLLEQINTSPDTWYAAID
						LANALFSIPVHKGYINSLALCH
						NVIWRELDCFSLPRDTTLVHYI
						DDIMLIGSIKFLGVQWCGACRD
						IPSKDPADPMVLEVSVADRDAV
						WSIWQALIDESQQRPLGFWSKS
						LPSSADNYSPFERQLLAYYWAL
						VETERSTMGHQVTMLPELPVM
						NWVLSDPSSHK/ANGLAGWSG
						TGKKHDWKIGDKEIWRRGMW
						MDLSEWSK/D/VKIFVSHVSAH
						QRVTSAEEEFNNQVDRMTRSM
		<u> </u>				DTTQPLYPTTPVIAQWAHE
4506	34874	Α	4547	1	1236	OFFICE OFFICE AND A PERSON WAY
4507	34875	Α	4548	1	1467	GEKGNDQTFRKLLDTGSELMLI
						PLRVVIPTTSLFNSPIWPVQKTD
						GSGRMRVDYHKLNQVMTPTA
						AAVPDVVSLFEPINTFLGTWYA
	1					AIDLANALFSIPVCKAHQKQFA
						FSWQGQQYTFTVLPQRYINCLA
	}					LCHNLIQRDLDHFLLPQGITLV
	1					HYIDSGPFIK*PEAASFEWGPEQ
	1					EKALQQVQAAVQAALSIGPYD
						PADPMVLEVSVADGDAVWSL
						WQAPKGESQWRPLGFWSKALP
						SSTDNYSSSDVQLYTDSWAVA
						SSLAG*SGTWKKHDWKIGDKEI
						WGRGMWMDLSEWSKTGKIFV
						SHVNAHQLVTSAEEDFNNQVD
						RMTRSVDTTQPLSPATPVVAQ
						WAHEQSGHGGRNEGYAWTQQ
						HGLPLTKADLTTATAECPICQQ
	1					QRPTLRPRYGTTSQGDQPATC
						WQVDYIEPLPSWKRQRFLLTGI
						NTHSGYGFAYPPCNASAKTTIH
						GLIACLIHCHGIPHSIASLYRER
			<u> </u>	L		GTHFTDKEVQQWAHAH

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ 1D NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4508	34876	A	4549		1602	NLSPILPQDLWPFTRVTVHWGK GNDQTFQGLQDTGSELMLIPGD PKRHCSPPVKVGSYGGQVINGV LAQVRLTVGTVGPRTHPVVISP VPECIIDIDILNSWQNPHIDSLTG RVKAIMVGKAKWKPFEPLLPIK IVNQKQYRIPGGIAEISATIKDL KDAGVVIPITLPFNSPFWPVKKT DGSWRMKVVYCKLNQVVTPIT AAVPDV/VVSLLEQINTSPGTW YAAIDLANAIFSIPVHKAHQKQ FAFSWQGHQNTFTVFTILLHIH KVGHAQQHSIIKWKWYIHDGA RAGSEGTSKLNEEVPQMPMVT TSAALPSLPRPAPMASWGVLY DQLTEEEKTRAWFTDGSARYA GTTQKWTAAALQPLSRTSLKG SGEGKSSQWAELQAVHLVVHF SWKDKWPDVRLYIDSWAVAN GLAGWSGTWKKHDWKIGDKEI WGRGMWMDLSEWPKPVKIFG SHVSAHQWVISAEEDFNNQVD KMTCSVDITQPLSPATPVITQW AHKQSGHGGRDGGYTWAQQH GLPLTKTGLAMATAECPI

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4509	34877	A	4550		1891	MLSSTQNAGGSYQRVRGALDT QEWKWGEVSPRTLNVDGRAL VSVANTHGTDRPAYTLNPQSR DQRSGVITLGYKRPLEREDLFE LKESDSFCTACPIFEKQWRKEV LRNQERQKVKALNKLDEALCP GIILTQSTDSNANLFQKQPHRHT QTSGRWQIIIFCEHSSDFGWNG YGYAVALLVVVFLQTLILQQY QRFNMLTSAKVKTAVNGLIYK KLGWSGKVSWLILHDVGHGIM EGYIAWGKGSDVRITWEKKST EMRTRPAQKMALLLSNVSRQK FSTGEIINLMSATHGLDSKPQSP LVCPFSNPNGRISPLARAGLAD HYRVTHLQILKLYAWEPSYKN KIIKIRDQELEFQKSARYLTVFS MLTLTCIPFLTKISLGRLEDFLN TEELLPQSIETNYTGDHAIGFTD ASFSWDKTGMPVLKESIRIRIEQ VLNQLSLFETVDYPGSVAYVSQ QAWIQNCILQENILFGSIMKKEF YEQVLEACALLPDLEQLPKGD QTEIGERAVNISGGQQHRVSLA RAVYSGADVYLLDDPLSAIDV HVGKQLFEKVIGSLGLLKNRTH ILVTHNLTLLPQMNLIVVMKSG RIAQMGIYQELLCKTKNLTN\FT
4510	34878	Α	4551	2	542	KSSVNKKKVGEWEESGRGS LTSAKVKTAVNGLIYKKVSLAT LCVYFLLDERIILTAPKVFTSMS
						LFNILRIPLFELPSVISAVVQTKI SLGRLEDFLNTEELLPQSIETNY TGDHAIGFTDASFSWDKTGMP VL/NRGSEAYVSQQAWIQNCIL QENILFGSIMKKEFYEQVLEAC ALLPDLEQLPKGDQTEIGERVR
4511	34879	Α	4552	1	667	IETNYTGDHAIGFTDASFSWDK TGMPVLKESSVAYVSQQAWIQ NCILQENILFGSIMKKEFYEQVL EACALLPDLEQLPKGDQTEIGE R/GKETAVNISGGQQHRVSLAR AVYSGADVYLLDDPLSAIDVH VGKQLFEKVIGSLGLLKNRTSH SVCHYTLLAVPHLLEVQILTGN FIQSLGFNYHEYANNSNAYIVN LDLFPGFQTCVYKLLSPIRCLIC
4512	34880	Α	4553	201	336	QQTPGKAVHAPFIADQSLT*EL VSVFPQFQLFPYRR*DSHSGKS
4513	34881	Α	4554	3	515	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4514	34882	I IA	4555	<u> </u> 	852	MPCTTSGLDKVPTSKKALTRY
		` `				GYGSLTLDFSGSIIPCKMCLSPS
						TRIVRPSQPRGTEDPDETGDTEF
		İ				VNSDESFLLEGTASPSPVVAVSP
						PRPMLPSAFPPLSEDINPVLPEA
		ŀ			,	TVLASPEVVAKQTHVDSPRKPL
		į .				STFLFASRPVTKLKSRQTPGGE
		l				VDSVTCEEKTDGPWRKTVDYC
						KLNQVVTPIAALVPDMVSLLV
			İ			QINTSSDTSYAAIDLAKAFFSIP
						VYKAHQKQFIFSWQAQQYTFT
						VLPQGYIISPALCHNLIRR/DLD
		ĺ			Ì.	HFLLPQDITLVHYIDDIRL
4515	34883	В	4556	288	327	THE EDI QUITE VITTIBBILE
4516	34884	A	4557	51	598	LFGGCHTSGGLAVRVPRMPRG
	310					SRSRTSRMAPS\ASRAPLK*ELE
		ļ				PRQAQVAQPPA\AAPPSAVGSS\
						AAAPRQPG/LFMAQMATTAAG
	1					VA\VGFCCGGHTLGHGI\TGGLS
						VGGKLIA*ALRRP*HQFNQGSF
						RGTQAKHSKQQPALPLLWRIKT
,						SFREVVPPEPRVTIQGFCGGFPM
<u> </u>						RLLETVPDL
4517	34885	Α	4558	1	10434	MTVIRSGIAYILHLKSYDVNIQT
	1					GSNACNQPTHPNGDCSHFCFPV
						PNFQRVCGCPYGMRLASNHLT
	+					CEGDPTNEPPTEQCGLFSFPCK
	-					NGRCVPNYYLCDGVDDCHDNS
	1					DEQLCGTLNNTCSSSAFTCGHG
						ECIPAHWRCDKRNDCVDGSDE
1				ŀ		HNCPTHAPASCLDTQYTCDNH
	1					QCISKNWVCDTDNDCGDGSDE
						KNCILNCTASQFKCASGDKCIG
						VTNRCDGVFDCSDNSDEAGCP
						TRPPGMCHSDEFQCQEDG
4518	34886	Α	4559	24	849	ATGRCCCGLAPGFPLCWVLYP
						GGRGSA\CPEPHVLRTGSPLQRE
				1		QRTNGRTDLSSLLPNLNFDSPP
						RCKHKNQLAITLRKRIRKLATS
						LFSSTIFRISGTSVIISAPGAGLPL
						PALFPTRCQPKFSRSIDPTGKAV
		-				QTADIRLSARATLWLGGSIEESP
		1		1		VLCSTLRLLLRRLPPPLTWTSPN
	<u> </u>					RPTQPCTAQTQTNQSVGIAAPS
				1		AIRVIYPESVVLNAVIYLPGDPE
						VSGLPRAFKRRFSVEVRLDCGT
						FKLLLVYCTHPGDKKVNTCKT
				<u> </u>		GALVAF
4519	34887	С	4560	192	449	
4520	34888	Α	4561	1	786	
4521	34889	Α	4562	3	14073	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4522	34890	В	4563	58	1282	
4523	34891	В	4564	1	684	
4524	34892	Α	4565	1	1356	MGRKYTLKWEFEEGFTEKKEL
						KKVTSEGYITPVEIYDYRQYCY
						ALQRPIATTQIDDVRDGHTTRL
						AKLEKQEQTHSKASRRQEIIKIR
						AEPKEIETQKTLQKINESRSWFF
						EKINKIDRLLARLIKKKREKNQI
						DAIKNDKGDITTNPTEIQTTIRE
						YYKHLYANKLENLEEMDKFLH
						TYILPRLNQEEVESLNRPITGSEI
:				ļ		EAIINSLPTKKSPGPDGFTADFY
						Q\MLEVLARAIMQEKEIKGIQL
						GKEEVKLSLFADYMIVYLENAI
						ISAQNLLKLI/SNFSKVSGYKINV
						QKSQAFLYINNRQTESQIMSER
						PFTIASKRIKYLGIQLTRDVKNL
						FKENYKPLLNEIKEDTNKWKKI
		ŀ				PCSWVGRINIVKMAILPKVIYRF
				:		NAIPIKLPMTFFTELEKTTLKFI
						WNQKRACITKSILSQKNKAGGI
			:			MLPEFKLY/YQGSSTQTAWYW
4525	34893	Α	4566	1	1102	MANCDINRKDEKGGKEKKDRS
						KSKSLMDTLKRQLSAKQKPKG
						KAGKPSGSSADEDTFSSSSAPIV
						FKAVRAQRPIR/STSLRSHHCSP
						MPWPLRPTNSEETCIKME\PSPP
						LNGVRKDFHDLQSETACQEQA
						NSLKSSASQNGDLYLRLDEHVP
						VVIGLLPQDYIQYTVPLDEGMC
				İ		PLEGSSSYCLDSSSTMEVSVVPS
		ŀ				QVGGRSFPEDESQADQNLVVA
						PEIFVDQSMNGLLTGTTGVMLQ
		1				SPRVGPHHVPPLSPLLPPMQNN
						QIQRNFSGLTGTEAHMAESMLC
						HLNFDFNSAPGVARVYVSVQSS
						GPMVVTSLTEELKR\LAKQGWL
						WPPLKSVRRCVLARRSLYTKQL
4506	24004		15.60	264	((1	NQEEGTELNLGSSCLLC
4526	34894	Α	4567	364	661	PFHFTCFSCKVYFADPGSAARS
						VPGSPSAVCAQCILCTGHCAVC
						PGLGEHHSSGRTLMKTKLHSK
						KLKPCYLLC*SKN*KTQGGSPK
4507	24905		1560	52	470	S*NVNKYLVTLI CISIIILPGPSAKTLSPVLSLSSPY
4527	34895	A	4568	53	470	
						TASFQPTFVRTFSHQTTYLSLGS
						VPVAQLKCSAGQQRGELLCRR
						GVWGSWISVSHFTEIATLPAAC
1						LEDGE\DFNLGGILDSSKYL*SIQ
L		<u> </u>	_	<u> </u>		KTNTHRIVDGKVVSETNITDVL

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO:	Nucleotide location of first	*	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO.	sequence	, nou	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4528	34896	Α	4569	1	1635	MGWTCKILFLVAAATGAHFLV
						QLVQSGAEVKKPGASVKVSCK
İ						ASGYTFTYCYLHWREPVSFICG
		1				RCILLTLLLGDCLWGEGAWTQ
į						GDVLQPSDRASFLAMGVNTTG
						QQVGDLSGDFPNSVGKACKCR
						EFHTLTPLAHTSSTTHETFPGMS
						HIALELSQGSSLLQCLEAQTQG
İ	1	į				QRQELTVSANEQPESRGHGCVL
						LCETQSEGKSVRAQAQTSLKGS
						QKRLGGARTLCTGLSPGQRKQ
						ERSKIDTLTSQLKELEKQEQTYS
						KASRRQEITKIRAELKEIETQKT
						LQKINESRSWFFEKIYKIDRPLA
				ļ		RLIKKKREKNQIDAIKNGKGDI
						TTDPTEIQTTLRQYYKHLYANK
						LENLEGMDKFLDAYTLPRLNQ
						EEVESLNRPITGPEIEAIIN/STPT
						KKSPGPDRFTAEFYQ\RSDVLA
						RAIRQKKEIKCIQLGKEEVKLSL
						FADDMIVYLEIPIISAQNLLKLIS
						NFSKVSGYKISVQKSQAFLYTN
1		1				NRQTESQIMSELPFTIASKRIKY
						LGIQLTKDVKDLFKDNCKPLLN EIKEDTNKWKNIPCS
4529	34897	В	4570	1	429	EIREDINKWRNIFCS
4530	34898	A	4571	1	897	MDLNYTLEQMDLTDIYRTFHPT
	3 1070	1.		•		TTEYTFYSTGHGTFSKTDDVIG
		:				HKMSLDKFKKIEMISNTVSDHS
						GIKLEINSERNLENHANTWKLN
				·		NLLLNECWVKNKMKMEIKKLF
					·	ELNDNNDTTYHNLWDRAKVVI
						RGKCIALNTYIKKSERAQTDNL
						RIKNKNHMIISIDAEKAFDKIQH
	•	ŀ				PFMIKTLSKISIRGTYLNLIKDIY
ŀ		ŀ				DKPTANIMLNGEKLKAFTLRTG
						T\RMNQGCPLPSLLFNI/VLEVL
ŀ						ARAIRQEKEIKGIQIGKEEVKVS
		İ				LFADYVIVYFENPTDSSRKLLEL
Ì	1			<u> </u>		IKEFSSFWIQD
4531	34899	Α	4572	1	1461	
4532	34900	A	4573	49	365	West Brown general age
4533	34901	Α	4574	45	534	VCHLEPGERCGPSRGCRAVGV
		1				QTEKMQTAGALFISPALIRCCT
						RGLIRPVSASFLNSPVNSSKQPS
						YSNFPLQVARREFQTSVVSRDI
						DTAAKFIG\AGSATVGVADSGA
						GIGAVFGSLIIVYARKLSLKQQL
						LFYAILGF\ALSEGM\GLFCLMV
<u></u>	<u></u>	<u> </u>	İ	<u> </u>	<u></u>	AFLILFAM

SEQ ID	SEQ ID NO	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4534	34902	A	4575	98	624	DWYSRSHPKELWEGSKKKSIN HSACSLQAHGGPISTSLNPHSKP RGRVSPPPGKRQQECRARPGRS PELAGPPPANVQETSQKNACAS RLSEPPGEGP\EPAAHPQPHRGS SSGPCSRGGYRQPLFPGPAASG VPASGSV/RSRIPGAPQGVALAR RGPQGSPSPAPRFFPATERQS
4535	34903	В	4576	1	604	
4536	34904	A	4577	3	331	LAPAPSAAWRTGLKALTPSST WMLCASE\HHVGSGCVGDHLA GCRQEKTLPCQR\YCVFCRRR ARSLQAQCGFSLTPALELLPVPF LKLLCPGPPRRRRICRILPGAGL
4537	34905	A	4578	1	871	
4538	34906	A	4579	3	510	GPPSRVDDFVAAAAAAVAPVV LYACPRHSPIPPWSIRGRRVVVT GFGPFGEHTVNANWIAVQELE KLGLGDSVDLHVYEIPVEYQTV QRLIPALWEKHSPQL\VVHVGV SGMATTVTLEKCGHNKGYKGL DNCRFCPGSQCCVEDGPESIDSI IDMDAVCKRVTTLGQCI
4539	34907	A	4580	1	285	MAPGALPALGEEEGPGASGLSA ELGHLSAGSRAFRETSVDSALD TPFPAGTFVRLEFKLRQTE\SGR RKDWKKPKCKVQPERRKQKCL TCVKLEC
4540	34908	В	4581	1	228	
4541	34909	A	4582		697	MGLERPVDRVMWLPGALWNS AVVSAPVGEEWALAGTGNQGL QDIQGMHCPEEGISQIHGRDHR NAKDSHTGVWCSCTLGIISTIIIR PKCRFSIDRSDSDYLPTSSCRRD PGGAEPCQDRPRVEQLCSVLAN RSGPLAKCHWYESPVSYTQVC VSDLCQYGTGNRMLCTMLEAY VQLCALRCALPARVASQPGMQ LRVACPANSYYDSCGPPFPATC ASLNSSAPCTLQCTVSCFCLEGF ALEAG\SSVPHACCGCHLQGRY I/APGPWPSATGMRAPCPTRRC VSLTSASMARATACCAPCWRP TSNSAPCAARCLPAWRASLGCS YVWRVQPTATMTPVGHPSRPP VLASTPPRPAPSSAQ
4542	34910	В	4583	1	208	
4543	34911	Α	4584	2	230	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
4544	34912	ĪA	4585	106	669	GCGCVLLPGGVGAPGHSPEGP
		`				VPPRDARQLRTCGLAGTLHLST
						SCGHPTGAWRGALGSQWNGH
						NPVQDHLQEAQPWFLEFDRG*
						RCFWRMATSFPRYPTWDDY*R
						LPCCRCLPQCKKPAETTGCLPIS
						GEKTHGGVGDLLGEAPARLRQ
		l				WASQRQPATDLPA*ASRGRPA*
						GKDTHRACVAGEAAQDA
4545	34913	A	4586	2	597	TPKGGIRLGLAKLGCPTAWINP
1373	34913	^	17300	2		YGRGMPLAHSVLSSGARVLVV
						DPDLRESLEEILPKLQAENIRCF
		1				YLSHTSPTPGVGALGAALDAAP
						SHPVPADLRAGITWRSPALFIYT
						SGTTGLPKPAILTHERVLQMSK
					,	MLALSGATADDVV\YTEVLPLY
						HVMGLVVGILGCLDLGTSLAS
						YGLRVYFILWSVLGPSRRTLCL
4546	34914	A	4587	9	573	EEEERKKKKK\KKEEEEGEEE
14340	37717		1307		373	GGGMGEKKKKEEEEEGEEGE
						KEREKER\EKERKKKKEERKEK
						ERERKKKRERKKEKERERKEG
		İ				EGERERKSTECTSSSYI/IKKLVV
						KQPQAAPSGEIPEEGIAVLGGDS
						SMPVIVPEDLPVGQDVEVEDSD
						INDPDSLILVSSQAGGGGVITAY
						CNLEHLGSSDPPT
4547	34915	A	4588	1	297	CIVELITE GSSBITT
4548	34916	A	4589	114	752	DGSAAPRATSDSFTYTVCVSEF
						PVDDFMELGRSIPDTQL/DAVIE
		Ì				SQKANQCAVLIYT/SGTTGIPKG
		ļ				VMLSHDNITWIAGAVTKDFK/P
						TDKHETVVSYLPLSHI/AAQMM
						DIWVPIKIGAL/IYFAQADALKV
						RLSKDLGSDFILLGSPVGLRPST
						KRLPVLSKLGHTYRRVVWVEE
						SSGPHTISNQNNYRLQGPMMK
						LKRHFVAQKYKKQIDHMYH
4549	34917	A	4590	1	837	MVTQKLPNAQENLKHAERQAA
						GCCPGRSHIFQHVGPGASESLR
						GEGCSTHPEAQGAQERCEQWK
						KDQHWCLASHTDVTQQWGRH
						IVQEGGTHRGPSAVLSLRTALD
						EG/ARGGCSHPITAQLPLQLRHL
						PRPPPAPAR\PSPAPPAATSPPT\P
					1	PPAPAR\SSPAPPAATSPPTASSG
						ACAALPQLPLQLRHLPRPPPAP
1						AR\SSPAPPAAMSPP\RPPAPARP
			1			RLRRSTACATALRPGERGSAAA
						QPGARSETSFCRLG/AAAAVLD
1						PAFISSQALACPVVGVI
L		L	I	L	1	T. T. 100 AUTUCE A A O A E

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide inscrtion)
4550	34918	Α	4591	3	377	
4551	34919	В	4592	1	1632	
4552	34920	Λ	4593	1	1224	MDGTNRFYLYVWETERQQDV
						EHVARCILCYDKCRPDPECPAG
						TPGPQEVDVDLVFVVDSSYGV
						DADVYRGSLSLADAALEDLEV
						AEQPGASHRGARVALVTHTTP
						NF\GRGFHLTTYGNRKQMQRH
						VREASARPLQGTAPPGHALEW
						TLENVLLAAPRPRKAQVLFAIV
						ASETSSWDREKLWTLSLEAKC
						KGITLFVLALGPGVGTHELAEL
		1				AELVSAPSEQHLLRLQGVSEPE
	-					VNYAQGFTRAFLNLLKSEQSPG
						TGAPWVEWGEGFTEPGIWACR
						WTNQYPPPELTEECGGLHRGD
						TVLQLVTPVNRFMYAAKENSL
						KRKTKANFHLAVELELDESYFR
		1				AYYEGTLYEVSALPLQRSNELL
						QKWSLFHGSNGRRVSGSHPEV
		<u> </u>				ALQQGGTGLPAVLVWQLWRQ
4553	34921	Α	4594	266	556	HKVQQICYRLRLVSQILFSINQT
						LAERQIVTFTVYPDTERDRETR
						NLADLKQIKIDLGKFSDNPDGY
						IDILRGLRQSFDLTW\RDIMLLL
						NQTLAPN
4554	34922	В	4595	1	735	
4555	34923	Α	4596	70	624	PTAMVEEGIAAGGVMDVNTAV
						QEVLKTALIHDGLARGIREAAK
					,	ALDKYVYQSQYCGFLQPDQKL
						ATQGKKGMGVHGVKRKSS*M
						ASVLPGNLRKRRQAH\LCVLAS
						NCDEPMYVKLVEALCAEHQIN
						LIKVDDNKKLGEWVGL\CKIDR
		1				EGKPRKVVGCSCVVVKDYGKE
1556	2.402.4	 	4505	142	600	SQAKDVIEEYFKCKK
4556	34924	Α	4597	145	682	SWRNRTVSNGSAVSASSVHLCF
						AECKALCGERILTDGSDVSRPTI
						AAGGVTDVNTALQEV\LKTALI
						HDGLAR\GISRTWPKAL\DKRQ
						AH\LCVLASNC\DEPMYV\KL\V
	1					EALCAEHQINLI\KV\DDNQET*
						EKWV\GLCKI\DREG\KPRE*WL
						VGS*CSSLRTIGKESQAKDVIEE YFKCKK
4557	34925	A	4598	252	590	RSLDLVWWQLSGGLAGSAKPK
4337	34723	^	4370	232	390	PCTPVKQSTVMSFPSHKEQYFL
						MDGKKK/YDKESEKYYSILEKH
						LNLSAKKKESHLQENSSGPSVS
						TKLINLFSKRLLCAFLPAQLTPY
		1	1	1	ļ	SFCS

SEQ ID	SEQ ID NO:		SEQ ID NO:	ł .		Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4558	34926	I A	4599	1	1662	
4559	34920	A	4600	4430	4904	LIKGTRCLGRRPSFAYSRSNI*A
4339	34921	^	14600	14730	4904	*S*P/EGTSAQLAELLALTLALE
						LGNGKRINVYTDSKYAYLILHD
						HAAIWKERAFLTSGGTPIKYHK
						EIMELLHTVQKPKEVAVLHCQS
						HQKESSPLEDTTTAGPLLHPYP
						AGSSPERSSPSSQQLLGDLFRGII
4560	34928	A	4601	1	2630	MEQANHPVRLINVVCKDTLKK
4300	34720	A	4001	'	2030	VQQETSCPLTHVHYAEAITGRC
						TAPEDKGSLDQKPPTDDPTGCP
				ļ		WQVPAHVITLTETWVCLTIEGQ
	:					EIDFLLDTGVQKPNGQWRLVQ
						DLIPIKEAVIPLYPVVPNPYTLIS
				•		QIPEKAEWFMALDLKDAFFCIS
						LHSDSQFLFAFEDPTNHTSKITR
						TVLPQGFRDSPPLFGQALAQDL
						GHFSSPGTLVFQYVDDLILATSS
						EASCQQATLDLLNFLANQGKV
						VPNLWGKLPLNTTRKSWSYCT
!						QCKNPRRWQSYTAKAIKKQLA
						EAGPVTAILLLLIFGPCIFNLLIK
						FVSSRIEAIMLQMVLQMEPQMS
						STNNFYQGPLDRCTDPLSGLES
						SPRCSEAPCLMSQWTGDIEYDL
				ŀ		LLPPIPHQTTLCDLQNLKGIFSR
		Ì				YHRKWYGEILALLTPTANVCG
						HSQVPHACSIYHDPVTWNPQG
						LLPKSLYGVTKWGDKEHFEWG
	į					SQQQRAFYELK\KKLMSAPALG
	ļ					LPDLTKLFTLHVSDREKKMAV
	ļ					RVLTQTMGPWLGPVAYLSKQL
				İ		DGVSKSWPPCLRALAATALLA
						REVDKLTLGQNLNIKAPHAVV
	ļ					TLMNTKGHHWLMNARITRYQS
						LLCDKPHITIEVCNTLNPTTLLL
						VSESPVEHNCVEVLDSVYSSRP
			İ			NLRDHPWTSVDWELYVDGSSF
						INPQGESVWGIIQGKRPIKLWG
						KRRKVSARDLAIIGGSVEAPKL
4561	34929	A	4602	11	506	FLALTSRFLFVLLNEETRSHLEK
		1		1	-	SLCWKVSPHIKMDLLQWIQSK
						AQSDGSTLQQGSLEFFSCLYEIQ
						EEEFIQQALSHFQVIVVSNIASK
						MEHMV\SSFCLKRCRSAQVLHL
	}					YGATYSADGEDRARCSAGAHT
						LLVQLPERTVLLDAYSEHLAAA
				}		LCTNPNLIELSLYR
4562	34930	A	4603	3	381	
4563	34931	A	4604	3	483	
4564	34932	A	4605	3	410	

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO:	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
110.	sequence	Hou	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4565	34933	A	4606	2	249	SADAPMFDMGVNHEKYDNSL NII/SVMKAGPVEKRPAWHPMD TLP*LAPRSLFLCSNASCTTNCL EPLAKVIHDNFGIVEGLMV
4566	34934	A	4607	2	481	LAPLVKEIHDNFGMGEGLMTT GHAITATHKTADGPSGKLWRD\ GRGAHQNIIPASTGAAKAVGK VMPELNGKLTGVAFRVPTANV SVVDLTCRRQKPAKYDDIGKV VRQAPEGPLKGILGYTEHQVVS SDFNSDTHSSTFDAGAGIALND HFATLSPPPH
4567	34935	В	4608	79	278	
4568	34936	A	4609	2	1201	PSTACRNSARACSTVSRIFFCVA SRATSLRTPMGKVKVGVNGFG RIGRLVTRAAFNSGKVDIVAIN DPFIDLNYMVYMFQYDSTHGK FHGTVKAENGKLVINGNPITIFQ ER\YPSKINWG\DAGAEYVLEST GAFTTMENAGAHLQGGAKRVI ISAPSA\D\APMFVMGVNHEKY DNSLKIISNA\SCTTNCL\APL\A KVIHDNFG\IV\EGLMTTVH\AIT\ ATQKTV\DGPSGK\LWALMGPR GFFQEHQSLPFTGGC/ARVVGQ GSSPELERGKLTWAWAFRCPQ LPKRVNGWDL\TCRL\EKPCPK YD*HQGRVVKAGRRKGPLQGA ILGLQLSNPGGSPSGLSTSDNPL LPPFDAWGLAFALQRTHFCSKL IFLGIDNGILGYSNQGGWDLHG PPWPTWAFQGS
4569	34937	A	4610	61	226	WRIMPTKKVMITMGRTTQRRM LES/SQQFWPCHLH*KLVPSCLQ LGCLVFHFRRER
4570	34938	A	4611	153	495	QHAAECKAHAGLPGLPLPARK LASRHGAPRWRQSGVGPGGKV ENYGRRRL\PGTRHPQSLSHKP AKKIDVARVTFDLYKLNPQDFI GCLNMKATFYDTYSLSYDLHC CGAKRIMK
4571	34939	Α	4612	1	643	
4572	34940	A	4613	286	698	ESDNNLTQGTSI*QGTRHPQSLF PLSPAKKI*CGPVLTFLTCYKLN PQGLSLGCLNIEGRFFMDYVIPF PIDLALLLGAKRIMKG\TLHWA LFSMQTTGPRA/VFTSCYLQQL LDATEDGHPPKGKASSLIPTCL KILQ
4573	34941	Α	4614	59	294	
4574	34942	Α	4615	1	2253	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4575	34943	A	4616	12	197	LARSGMGFYRRGLLGEVKGRV
,		'				EGNRMWHVIVRTSPNHRYTFT
						LKTHPSVVPGSIAFSLPQ\RPWS
4576	34944	A	4617	302	441	
4577	34945	A	4618	944	1257	RLPFSPRSVGPTPQAPRLLCNG
		İ				WRQLPTTFFTELEKTTLKFIWN
		l				QKRALIAKTI/LKPKNKAGGITL
						PDFKLYYKATVTKTARFLLYK
				:		VSQIDNTDLFDPVKIKFE
4578	34946	Α	4619	1	1370	MNIDAKILNKILANRIQQHIKKL
}			ļ			IHHDQVGFIPGMQGWFNIRKSM
						NVIQHINRTKDKNHMIISIDAEK
						AFDKIQQPFMLNTLNKLGIDGM
						YLKIIRAIYDKPTANIILNGQKL
						EAFPSKHGTRQGCPLSPLLFNIV
						LEVLARAIRQEKEIKGIQLGKEE
		Ì				VKLSLFADDMIIYLENPIVSAQN
1						LLKLIGNFSKVSEYKINVQKSQ
						AFLYTNNRQTESQIMSELPFTIA
						SIRIKYLGI*\LTRGVKDLFRENY
						KPLLNKIKEDTNKWKNIPCSWV
•		İ				GRN\NIMKMAIL\PKVIYRFNAIP
•					,	IKLPMTFFTELEK\TTLKFIWNQ
		Ì				KRARIAKSILRQKNKAGGITLP
ŀ						DFKLYYKATVTKGAWFQHHK
						HTLIKEPLLDSVFSPNRPDHGK
						KQDKQPQTKNIANASADSKNT
						QQMNGFVTGAATSFIPKDRTAS
		l			•	SLCGCTGRRRQSVAKYLRIRPHI
						NVPSFTYYK
4579	34947	Α	4620	2	671	WHQNLALTRASGSFHS/WEEG
		l				KGGADMSHEICVANLQVYVRS
						TDFDRTLMSAEANLAGLFPPNE
						VQHFNPNISWQPIPVHTVPITED
						RLLKFPLGPCPRYEQLQNETRQ
						TPEYQNRSIQNAQFLNMVANET
				1		GLTNVTLETIWNVYDTLSCEAP
						SPPWGRKPPLERLWPRPRELTC
						PLRYT\QTHGLLLPPWASPQTV
				}		QRLSQLKDFSFLFLFGIHEQVQ
						KARLQG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
4580	34948	A	4621	11	1923	DSVAFEDVAVNFTQEEWALLG
-300	34540	``	17021	1	1,523	PCQKNLYEDVMQETIRNLDCG
						HETEFVEYENLRDPMNMPLHT
						TDGPHKCKICGKGFDCPSSLKS
į						HERTHTGEKLYECKQCGKALS
						HSSSFRRHMTMHTGDGPHKCK
				İ		ICGKAFVYPSVFQRHEKTHTAE
						KPYKCKQCGKAYRISSSLRRHE
						TTHTGEKPYKCKCGKAFIDFYS
						FQNHKTTHAGEKPYECKECGK
						AFSCFQYLSQHRRTHTGEKPYE
Į.						CNTCKKAFSHFGNLKVHERIHS
						GEKPYECKECGKAFSWLTCFLR
						HERIHMREKPYECQQCGKAFT
						HSRFLQGHERIHTGEKPYECKE
						CRKAFSWLTCLLQHERIHTREK
-						PYEGKOCGKAFTHSRFLOGHE
						RTHTAKKLCECLSTVSARKSVD
1						LIIASYPLFLNLFSTPKTLRNCSY
						RRHERMHTGEKPYECKQCSKA
						LPDSSSYIRHERTHTGEKPYTCK
						QCGKAFSVSSSLRRHETTHSAE
						KPYECKQCGKTFHHLGSFQIHM
						KRHTGDRPHKCKICGKGFDRPT
		ŀ				LVRYHERISTGEKPHECKQCGK
			İ			AFDHLGSFQRHMIRHTRDGPH
						KCKICGKGFDCSSTLQSHERTH
						TGEKKLYECKQCGKALSHSSSF
						RRHMTMHTGDGPHK\CKICGK
4581	34949	Α	4622	11	256	MGKGSFKYAWALYKQKAECE
						RGVTIDTSLWKFETSKCYVTI\K
						DFIKNIITGTSQQGQTASVAAFC
		l				ILSSCPASWKNQVSHRLGG
4582	34950	Α	4623	173	717	SINAVASTRRTIEKFEKEAAEM
						GKGSFKYAWVLDKLKAERERG
1						ITIDISLWKFETSKYYVTIIDAPG
		1				HRDFIKNMITGTSQADCAVLIV
						AAG\VGEFEA\SICQNG\QTREH
						ALLAYT\LGVK\QLICRVNKMD
İ						STEPPYSQKRYEEIVKEVSTYIK
	Ī	ĺ				KIGYNPDTRAFVP\ISGLNGDH
	İ	1				MLEPKC
4583	34951	A	4624	3	525	GCPSGPGHRCVAGHGAPGAVC
"00"	ועקדען	()	1027	Γ		RHVPTAWPGYSRCSPGPGPRGV
					1	EAVGHQRHRAPETHSTPAADR
		1				HRRGLPGS\KSDSAMEPSPSPAP
	,					1
						QAQPPKPVPKPRTVFGGLSGPA
						TTQRPGLSPALGGPGVSRSPEPS
		1				PRPPPLPTSSSEQSSALNTVEMM
			<u> </u>	1	<u> </u>	PNSIYFGLDSRGRAQAAQDK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4584	34952	Α	4625	2	448	RRAHACARRRKKEMLGVNVL
						TSHSSQERMKLTFKKKAVNFA
		ľ				DAAAAQGPLLPAMVNPTMFFH
						IAVDGEPLGCVSFEVRGLESKK
						*LLI*SIKLC*QIG\LFADKVPKT
						AENFHALSTGEKGFGYKGSCFH
				l		RIIPGFTCQSGDFTRPTA
4585	34953	Α	4626	1	751	GTRDATAEENRVLLAMVNPTV
						FFDIAVDGEPLGRVSFEVRGLD
				1		TKK*LLI*SIKLC*QIGGSSIFITS
						D*KNSCLPLIVQQCLLFLRILP\L
						FADKVPKTAGV*FFFKQKIFRA
						LSTG\EKGFG\YKGFPAFHRIIPG
						FMCQGW*LSHRHNGTGWQVH
						LMGRNFEDE\NFILKAYGVLGS
						LSMAKCLDPTKIGSPVFPSCTA
ľ						KT\EWL\DGQALWCFGK\VKKG
						LNIVEAMERF\GSRNGKTS\KKI
						TIADCGQLE
4586	34954	Α	4627	3	615	PECIIGIDILSSGQNPHIGSLTGR
				i		VRAIMVGKAKRKPLELPLPRKI
						LNQKQYRIAGGIEEISATIKDLK
						DAGVVIPTTSLFDSPIWPVQKT
						DGSWRMTVDYRQINQVATPIA
						AAIPDVASLLKQINTSPDTW/PI
						RPPISNGD*GVSGR*ACCLEPLA
				i		GPHR*ITSEASRILEQGPAIFCR*
					1	LLSF*ETALGLLLGFGGN*TFDY
4587	34955	Α	4628	3	354	DSWA\VANGLA\GWS\GTWKK
				1		HDWKI\NDNEIWGK\SMWIDLS
						EWSKTVKIFVSHESAHHIT*KSS
						AEEDFNNQVDRMIHSVDTTRPL
						SPATPVIAQWTHEQSGHGGRD
		<u> </u>		<u></u> .		GGYTWAQQHGL

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO: in USSN	Nucleotide location of first		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4588	34956	A	4629	281	1529	VRVLSPVEKELKLWKNTHKLL
		l	ļ			SYPTVGAAVTQLQNLTAMGVI
						GSHGARGQVVALNRQRQGDL
						QPFTRVTVHWGKG/NMQIFGGL
						LDTGSELTLIPGDPKHHCGPPV
		1				KVGAYGGQVINGVLAQVQITV
						GPQTHPVVISPVPECIIGIDILSS
						WQNPHIGSLTGRVRAIMVGKA
						KWKPLELPLPRKIVNQKQYHIL
						GGTVEISATIKDLKDTEAVTPTT
1						SPFNSPIWPVQKTDGSWRMTV
						DYCKLNQVVTPIAAAVPDVVS
	•					LLEQINTSPGTWFEWSPK\KAL
1					,	QQVQAAVQAALPFGPYDPADP
						MVLEVSVADRDAIWSLWNAAI
1						GESQRRPLGFWSKALLSSADNY
						SPFERQLLASYWALVETERLTV
1						GHQVTLRPELPIMNWVLSDPSS
						HKVSGAQQRSIIKLKWYIHDW
4589	34957	Α	4630	453	719	ARGSKHTGLIAQWAHEQSGHG
1						GRAGGYAWAQQHGLPLTKAD
1						LP\AMATAECPICQQQRPTLSPR
						YGTIPW/WAWDAPGGRGCWRL
						QKAGE
4590	34958	С	4631	122	325	
4591	34959	Α	4632]]	346	MAGEKVEKPDTKEKKPEAKKA
						DAGGKVQEGTGRYSRSAMYSR
1		İ				KAMYKRKYSAAKSKIEKKKEK
						VLATVTKPVGGDKNGGTQVF\
						QIITYSSYTQKVQLPKSTLKQRQ
4592	34960	Α	4633	115	905	GPCPQGAL EAFQTLHFCCLGLRQGTKRMA
4392	34900	^	4033	113	903	GEKVEKPDTKEKKP\EAKKVDA
İ						\GGKVKKGNIKA\KKPKKG\RPH
						CSRNPVLCSEGFGRYSRS\AMY
						SRKAI\YQEGSTFSPLKSKVEKK
	į					KKEKVLATVTKPVGGDKNGGT
						RVVKLRKMPRYYPTEDVPRKL
						LSHGKK\PFSQ\HVRKLRASITP\
						GTIL\IILTGRHRGK\RVVFLK\QL
						AKLAYLLC*LGPLVLNRVPLRR
						THQKFCHLPLSTKIDISNVKIPK
						HLTDAYFKKKKLRKPRHQEGEI
4593	34961	A	4634	2		FVALAAVLCRQCLPRAWVCRR
7,773	J4701	Γ.	+00+	-	330	AGQGSGRHYRAAICAELKKPLT
					·	IEEVAP/DPVGPHEVRVDVHFC
						GVNFGDILICRDQYQERPHLPFT
					,	PGPVADSRKGLPIRSCCPPYNL
					·	WHCDFCS
		Ц				WITCDICS

SEQ ID	-	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4594	34962	Α	4635	1	556	MGLKLNTHMDQTRGSLSGDTL
		ļ	1			EGSPSKRAKIILKRGGFRGILGG
	ļ.					VGVQERRTGQRPALPCHDLRR
						AEVVPDHRRGGPRPVGPHEDL
						WQIPEKVSLQEAAVLPITYGTE
						SFALEHRARTQPGEIVLVTAAA
						GATGLAVMWQQISSGQGNIAA
			}			AGSDEKCKLAM\EKGAQSSVN
						YSQGSLKDSATDQ
4595	34963	Α	4636	1	142	
4596	34964	Α	4637	2	368	
4597	34965	Α	4638	2	504	HKVGHAQHHSFITWKWYTRD
						WSRAGPEGT\GVPYDQLTEEEK
						TRAWFTDGSARYAGTIRRWTA
1						AALQPLSRTSLKESGEGKSSQW
•			ļ			AEVRAVHLVVHFTWKEK*PDV
						RLYTDSWAVANGLAGWSGTW
1						KKHDWKIDDNEIWGRGMWIDL
						SEWSKTVKIFVSHESAHHIT
4598	34966	Α	4639	182	840	RTAVKGNLPTTPVI/SQWAHEQ
						SGHGGRDSGYTWAQQHGLLIT
		1				KADLAMTTAECLISQQQRPRLS
						LQYSSIPWGNQPATWWQIDYIR
						PLPSWKGQSQDSQDRSRNQGV
						KVKVAPLTITPSDTTAKFLLHV
						PAALHSAGVDVLVPEGGMLPP
						GGTTTIPLNWKLRLPPGHFGLLI
						PLSQQAKKRVTVLAGVIDLDC
						QDEISLLLYNRDAKELYRYTAH
4599	34967	Α	4640	3	283	SRVSCSPPPLSPPPPLSPPPLSPP
						PLLSPPPPPLSPPPSPPPPVSLPPP
						PPVFSFPSSCP/PPFPPPLLPPLPPP
					-	PPLSPPPPPPPWSPSPPI
4600	34968	Α	4641	1	531	MGSSHCTQPGMLAAAVGQAVP
						GTDTGADSVSLSASCCHQHAIH
						CAQAIRAKGHLQAHTELPSAPT
						QPPSCARQCPKSGGDLGGRELY
	1					NNHEIRSGKHIGVCISFANNRLF
						VVSIPKSKTKEQILEEFIKVTGS
						YWVLGLSFDFRKWPV/GQCCQ
						SFHGRLRATPARRAERRDARSF
						AL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4601	34969	Α	4642	115	2405	ATAEGSGSSGVKERGAGIRKAE
	1	-				RRRTEASGGGGGRGRRRSWRR
	İ					AGAEGVSEADARGRGGKGREG
	}					KGGSRGGARAHRERARRRVEL
	1					DRVCCQRRELRPPFYNSSTRAG
	1					HREQRARVSRNPIPSDRISPPQP
	İ	į				NGEISGNMATEHVNGNGTEEP
						MDTTSAVIHSENFQTLLDAGLP
						QKVAEKLDEIYVAGLVAHSDL
	1	ŀ				DERAIEALKEFNEDGALAVLQQ
						FKDSDLSHVQNKSAFLCGVMK
						TYRQREKQGTKVADSSKGPDE
						AKIKALLERTGYTLDVTTGQRK
						YGGPPPDSVYSGQQPSVGTEIF
				i i		VGKIPRDLFEDELVPLFEKAGPI
						WDLRLMMDPLTGLNRGYAFV
						TFCTKEAAQEAVKLYNNHEIRS
						GKHIGVCISVANN\RLFVGSIPK
						SKTKEQILEEFSKVTEGLTDVIL
		1				YHQPDDKKKNRGFCFLEYEDH
						KTAAQ\ARRRFN*VGKVQGFG
						GNVGTVEWADPIEDPDPEVMA
		1				KVK\VLFVRNLANTVTEEILEK
		1				AFSQF\GKLERVKKLKDYAFIHF
						DERDGAVKAMEEMNGKDLEG
						ENIEIVFAKPPDQKRKERKAQR
						QAAKNQMYDDYYYYGPPHMP
						PPTRGRGRGGRGGYGYPPDYY
			:			GYEDYYDYYGYDYHNYRGGY
		ľ				EDPYYGYEDFQVGARGRGGRG
						ARGAAPSRGRGAAPPRGRAGY
		<u> </u>				SQRGGPGSARGVRGARGGAQQ
4602	34970	A	4643	2	369	LOLL COCOTUDO A VIDOVIDED
4603	34971	Α	4644	1	1002	MNAGCGQTHDCAYRQKRPED
						VNEEGRLEQRNRKRQDEWGPR DKPASSGYKAGTLDVENWNRA
						GEGLKHAHQKGLKVDSSAFCT
						CSLIRTVLMPLSPYYSAGQQAE
						SKNLKESVVPPTASIENKKQER
						EDKNWPILPPPVAETSVPPPSVA
	ł					GIETPIQRILRSAAIAGEPSGPCA
						FPISVRPDSNNPQQFIHEHTPLEF
						KLLNELKTSVVNIGVQSPFTLG
						LPESAFGAMRLLPFDVKHWAR
						TCLSASAYLTWNLNGQEMCTD
						QVRQNRAAGHGDIAEDMLLGN
						GP/YFRPGTSNGTKRAWATIPEE
						GVPVQSFLPFMEGSQEPSAQFL
	[ARLREAV
4604	34972	B	4645	1	575	. M. C. C. C. C. C. C. C. C. C. C. C. C. C.
4604	34972	В	4645	1	575	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
4605	34973	A	4646	11	776	VVLMRNLMSGTIFSMLANISML
						GSLVIIIQYITQEIPDPSRVPLVA
				İ		SWKTYPLFFGTAIFSFESIGVDL
			Ì			PLENEMKNARHFPPILTLGMPI
		İ				VTTLDIGMAALGYLRFGD\DTK
			}			GSIILSLPICWYLHGLSGEGPSQ
		1				SFSERETKAQVIVPRSEVNVPRR
						PVSEHSGRGEQLCGLSCRLYQS
						VKLLYIAGILCTYALQFYVPAEI
		İ				IIPFAISRVSTRWALPLDLSIRLV
						MVCLTSAPMTRSTPFCKYSTRG
						RRRWLEIPV
4606	34974	A	4647	1	1294	MGKDFMSKTPKAMATKTKIDK
4000	34974	Ι^	4047	[1	1294	
						WDLIKRKSFCTAKETTIIVNRQP
ŀ						TEWEKIFATYSSHKELISRIYTE
		ļ				LKQIYKKKTNNPINKWAKDMN
						RHFSKEDIYASKKHMKKCSSSL
						AIREMQIKTTMRYHLTPVRMEII
		ł				KKSGNNRQPIVGPCDNSVILLY
						KILANRIQQHIKKLIHHDQVGFI
				ŀ		PGMQGWFNTCKSINVIQHINRT
						KDKNHMIISIDAEKAFDKIQQPF
			,			MLKTLNKLGIDGMYLKIIIAIYD
						RPTANIILNGQKLEAFPLKTGTR
						QGCPLSPLLFNIVLEVLARAIRQ
						EKEIKGSQLGKEEVNLSLFADN
						MIVYLENPIISAQNLPKLINNFS
						KVSVCKINVQKPQAFLYTNNR
		-				QTESQIMSELPFTIASKRIKYLVI
						QLTRDVKDLFKENYKPLLNE/I
		ĺ				K/EDTNKWKNIPCSWVGRINIV
		<u> </u>				KMAILSK
4607	34975	Α	4648	2	711	WNRRRPCIAKTIL/SQKNKAGGI
	1					TLPDFKVYCKSTVTKTAWYWY
						QNRHIDRWSRTETSEITPHIYNH
						QIVDKPHKNKQWGKDLLFSKW
						CWENWVAICRKLKLNLFLTRY
						TKINSRWIKDLHEKLKTIKTLEE
						NLGNTIQDIGIGKDFMTKMPKA
					-	IATKAKIDKWDLIKLKSFCTAK
						ETIIRVNKQPTEWENISAIYPSD
						RSLISRIYKE\LKNIYKRLDAVA
						HTCNPITLKGQGRWIT
4608	34976	Α	4649	1	576	
4609	34977	A	4650	1	771	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	1	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4610	34978	A	4651		887	FPEDSEPISISHGNYTKQYPVFV
						GHKPGRNTTQRHRLDIQMIMIM
						NGTLYIAARDHIYTVDIDTSHTE
						EIYCSKKTAWEILDRPMVATCR
						MKGKHKDE\CHNFIKVLLKKN
	1					DDALFVCGTNAFNPSCRNYKM
						DTLEPFGDEFSGMARCPYDAK
						HANVALFADGKLYSATVTDFL
						AIDAVIYRSLGESPTLRTVKHDS
						KWLKEPYFVQAVDYGDYIYFF
						FREIAVEYNTMGKLLGLHHELL
				1		RRTQDYGHKAGCPESCLLSVRR
						CPPPQSKAHRESVEELIKGCRR
]						HAGFCACGHITT
4611	34979	Α	4652	1	2890	MVLLKVDPGLWGSSLRVLLKA
						DPPYGTQAHAERHGRALAGGL
ļ						GVGEQSQSLDLLRMSHTYGAL
						FLPRAAVSSWCASVRIRKIKKSP
						LLDGAPLLYEPDTWLGKWSSS
	•					WTLVFTHPFSAALTHSALTARS
	1	ļ				DTGSLTLSPDGKLYSATVTDFL
	1					AIDAVIYRSLGESPTLRTVKHDS
	•					KWLKEPYFVQAVDYGDYIYFF
						FREIAVEYNTMGKVVFPRVAQ
						VCKNDMGGSQRVLEKQWTSFL
						KARLNCSVPGDSHFYFNI
4612	34980	Α	4653	1	480	MEGVEEKEKVPAVPETLKKKQ
						GNFAELKIK\PKMAFVLRIRGVS
i						GVSPKVRKVLQLLRLHQIFSGT
						FVKLN/KASVNMLRIVEPYIAW
						GYPNLKSVNELIYKRGYGKINK
						KRIALTDNTLIVPSLGKYGILCM
						EDLIHEIYTVGKHFKEANNLLW PFKLSSP
4613	34981	Α	4654	3	279	
4614	34982	В	4655	119	177	
4615	34983	Α	4656	157	359	HQRCRK**TKLEG*TCRKIEVN
				ļ.		TDYIKP*E*EFWMIYYISIYLTSI
1		1				CSSICNENILQLENREKSRNT

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	i .	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
		1		sequence		
4616	34984	Α	4657	65	900	GLAFPPILQKESPHPCLWILFTSP
						VALSTSRPPCVISSAALCPRRHG
						FLPPRLLAPLAAAFCPRPRCFLP
						LPPRLFAPAPVDFCPRRCGFLPL
						PPRVFAVSCPRRRGLLPPLLFAP
						SAAAFCGFLAPSPWLFAAFGTR
:						CRGFLGLLVPAAAAFFPRRLRF
		İ				FLPRLFSPAAAVTFCHPCFLPPP
:						PRLSVPAAASFCRRGFQQEIRTV
					•	EIRGSGTRVGLQHQTGHLPRGP
						SSLPAELHA*ARVSPLEASPPSC
		1				TR/WMQQPEDLGPALTLGRICG
						VGACWEL
4617	34985	Α	4658	927	1157	ARRSTAPVDCK*LQP*AQAAL
						WVLVWDHEAARQPEGVVQPA
						PGLWPSSRRWRKFQAPGSSGV
					1	CHPGALQACPEELS
4618	34986	Α	4659	1	563	MKLVAVFDKQDLHHGGDDISA
1						SSMHTQSPERFTSASELGTNNV
						SAFSV\YQAASEIEVTSSVLHAS
						SQKGLSSQHLGFGAPQAGGGSF
1						RHLAPQRKEVLEEYFKYDPEH
						KLIFRFVRTLFKAMRLTAEFAIE
						ITHGGRDVERATGLVNKIHRQG
						CGDSFCNIGGNAKPYVCCGKE
		ł				YVSSSKHQNGIAI
4619	34987	A	4660	1	681	MGKYAEALRSQQKAVLMSVR
						VMGIEHPNTIQENMHLALHCFT
						SRQLSLALSLLQGAHYLM/LLV
		İ				LGE/DHPEVA/LLDNIRRVLHRV
						MEYDLSLCFLDNALAVSTKYQ
						GPKALKVALGHHLITSVYESKA
		l				EF\RSALQHQKEGLAAH/TSLGE
						DQEKTKESSEYLKCLTQLAVAL
						RRAMHEIYRNGSSNNIPPLNFT
						APSMASVLEQFKGINGILFIPLS
						QKDLESLKAEVAQ
4620	34988	Α	4661	2	443	VWQSGGDSITSKTNIITCVNCY
						LYTCIDSSFNQYHSILIVRARQD
						IWLPVALHRPWESSPFIHVINNI
						LQKILKRSTQFIFTLIAIIMGLIA
						VTVIAATAGVALRQSIQTVHFV
				1		DKWQKNSTRMWNS/QRIDQKL
		$oldsymbol{ol}}}}}}}}}}}}}}}}}}$		1		ANQINDLRQTVIW
4621	34989	Α	4662	2	377	FAFTWTDPDTHQAQQITWALL
						PQGFADSPHYFSQAQISSSSITY
		1				LGIILHENTRALPADHV*LISQT
						PISSTKQQLLSFLGMVRYFCLWI
	}	1				PSFTILTKPLY*FTKANLADPTD
						PKSFPHSSFRSL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
1	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
4622	34990	Α	4663	1	1095	MAKAVLSKKNKTGSTTLPDFKI
						YYKDIVTEEPGIETLRHTSSTKF
						SKGLPSRESFSEKQNKTTTHLH
						PGEINSFIAHTKPAWRSLHTDA
						HEIWCRDSDQVNSPGRSIPCPPA
						LCSMRKIHLRPLVLRPASPRNIS
						PILNPLHLIAALLPNPKPPFRPPL
						VSPDLNPQVKDISTPSWATDHV
İ						HLTVSLKPYHPYPAQCQYPIPQ
	ŀ					HALKGLKPVITRLLQHGLLKPI
1	i					NSPYNSHILPVLKPDKPYRLVV
	ļ					QDLRLINHIVLLPIHPMVPNPYT
						LLSSIPASTTHYSVLDLKHAFFT
						IPLHPSSQPLFAFTWTDPDTHQA
						QQIT*AVQPQSFTDSPHYLNQA
						QISSSSVTYLGIILIKAHVLSLPIV
4623	34991	Α	4664	655	2417	KKRESMNIDAKILNKILANRIQ
1025	3.571	-				QHIKKLIHHDQVGFIPGRQGWF
		Ì		Ì		NICKSINVIQHINRAKDKNHMII
		<u> </u>				SIDAEKAFDKIQQLFMLKTLNK
	ŀ	}				LGIDGTYFKIIRAIYDKPTANIIL
	·					NGKKLEAFPLKTGTRQGCPLSP
		ŀ				LLFNIVLEVLARAIRQEKEIKGI
		1				QLGKEEVKLSLFADDMIVYLEN
ł						PIVSAQNLLKLISNFSKVSGYKI
						NVQKSQAFLYTNNRQTESQIMS
						ELPFTIASKRIKYLGIQLTRDVK
						DLFKENYKPLLKEIKEDANKW
ł						KNIPCSWVGRINIVKMAILPKVI
		ł				YRFNAIPIKLPMTFFTELEKTTL
						KFIWNQKRAHITKAILSQKNKA
						RGITLPDFKLYYKATVTKTAW
						YWYQNRDIDQWNRTQPSEITP
						HIYNYLIFDKPDKNKOWGKGS
1						LFNKWCWENWLAICRKLKLDP
ļ						FLTPYTKINSRWIKDLNVRPKT
						TKTLEENLGITIQDIGMGMDFM
						SKTPKAMATKDKIDKWDLIKL
1						KSFCTAKETTIRVNRQPTKWEK
						\FSQPTHLTKG*YPESTMNSNKF
						TRKKQTTPSKSGRR\NEQTLLK
						RGHLCSQKTHEKMLTITGHQR
						NANQNHNEIPSHTC
		1	<u> </u>	L	L	LIVITALII PILLO

SEQ ID			SEQ ID NO:	•		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN 09/540,217	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/340,217	sequence	or peptrae sequence	deterion, \-possible nucleonde insertion)
4624	34992	A	4665	1	234	MQNQEKFGELISEWAPYGRPRP
	1					QIAVGDHHISVYDGEVNWRMS
					:	SLQASAAGKDYANRGTEWGEE
	1					PSQMLRHELLSGEMCVSRNRG
						QPLANSERRSEALSPVTPKKLIP
						ADGHKSDPGRTWMKLETIILSK
						LTQEQKTKHHMFSLISG*FAEN
				:		DGFQLHPCPS\GSLLWPSAGISF
						FGVTGLRASDLLSLLARGCPRF
:						LDTHISPLSSS
4625	34993	В	4666	1	2553	
4626	34994	Α	4667	33	272	
4627	34995	A	4668	1	3045	
4628	34996	A	4669	1	334	
4629	34997	Α	4670	159	245	FPIVASPWLLC*LMSFAGTWVK
		-				LETIILSKLSHGQKTKHRMFSLI
İ						DTVVVGSDDFPLLRVPGCCASS
						LNAAHTSVNIKLELT
4630	34998	Α	4671	122	359	TANLKNGRKFL*PIHLTKG*YPE
			į			STKNVNKFTRKKQTTPSKSGQK
						I*TNTSQKKTFMQPTGT*KNAH
						HHWSSEKCKSKP
4631	34999	Α	4672	2	66	RLVYADTCFSTIKLKAEDASTS
		1				ENMRCLVFCACDSLLRMIVSSF
				:		IRVPTKDMYSSFFMAA*YLLQY
		<u> </u>				HQVKSRRCFYE
4632	35000	Α	4673	519	899	SALVCHTCSNWQVHLGDSVFY
			:			RSEEQPLEPLPFSYLSSLFPGLHP
						DPVSSGSQQPS*MPHTDASVTS
			:			SHGLGGLAGRNSCIYPCCAPAL
						CADYLWGSPDLFLLLSFQHKG
						NVGVGLAHSPQFQQGN
4633	35001	Α	4674	1	278	
4634	35002	Α	4675	158	592	GYYWRPSFQSLRENNECQRKS
						NSVNAGCLNCDHCVLGIYQQH
1				ŀ		*QNYFSFDIHYFLSETGRKVSSS
						I/AYFTIGETEALSGKVVPYWQQ
		-				AAGQGCTLHLLLPQTRLFPGKG
						RRQPGLLREF
4635	35003	Α	4676	302	721	
4636	35004	В	4677	1	871	
4637	35005	В	4678	1	559	

SEQ ID	_	1	SEQ ID NO:	l .	1	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		02/240,217	sequence	or peptide sequence	deterior, y possione nucleotide insertion,
4638	35006	Α	4679	3	2386	RVATMAPHRPAPALLCALSLAL
						CALSLPVRAATASRGASQAGAP
ŀ		ļ				QGRVPEARPNSMVVEHPEFLK
ŀ		ļ				AGKEPGLQIWRVEKFDLVPVPT
		ŀ				NLYGDFFTGDAYVILKTVQLRN
			:			GNLQYDLHYWLGNECSQDESG
						AAAIFTVQLDDYLNGRAVQHR
		ŀ				EVQGFELAP\FLGYFKSG\LK\Y
						KKGGVASGFKHVV\PNEVVVQ
						RLFQ\VKGRRVVRATEVPVSWE
						SFNNG\DCFIL\DLGN\NI\HQWC
						GS\NSNRYERLKATQVS\KGIRY
					·	NERSGRA\RVHVSEEGTEPEA\M
						LQVLGPR\VALPAGTEDTAKED
İ						AAN\RKLAKLYKVSNGAGTM\
						YVSLWAD\ENEFTQGA\LKSED
						CFILDHGKDGKIFVWK\GKHAN
						TEERKAALKTASGFHSPRWDY
		Ė				PKQIQVSVPFLEGG\ETPLFKQV
		ŀ		}		FKNWRDPDQTDGLGLSYLSSHI
					,	ANVERVPFDAATLHTSTAMAA
						QHGMDDDGTGQKQIWRIEGSN
						KVPVDPATYGQFYGGDSYIILY
						NYRHGGRQGQIIYNWQGAQST
				ļ		QDEVAASAILTAQLDEELGGTP
						VQSRVVQGKEPAHLMSLFGGK
						PMIIYKGGTSREGGQTAPASTR
						LFQVRANSAGATRAVEVLPKA
		1				GALNSNDAFVLKTPSAAYLWV
						GTGASEAEKTGAQELLRVLRA
						QPVQVAEGSEPDGFWEALGGK
						AAYRTSPRLKDKKMDAHPPRL
4639	35007	A	4680	1894	2161	MFGLPNARAATSTAPFASHSLC
				ļ		LCFRILLLLGPGINLANPRNHLV
						LHQKFSILGRHFSL\ATEEPCISL
						ALAPSKRWECNSSS*RYENN
4640	35008	A	4681	1	1803	
4641	35009	A	4682	1	501	MTFQCVVNTHYLTYPRPQRFL
						YLVVVRPSCASWIMFVLIDRGY
		1				VFSYFPQSYGGFGSRILSKPIEV
						QVGGRSVVAQMWSNKCGQYS
						DLASLGCISR/YSAGSVYYYPSY
						HHQH/NPVQVQKLQKELQRYL
		Ì				TR/KIGFEAVMRIRCTKAKPTRH
						RHYGELEISITIIRAIGK
4642	35010	A	4683	350	623	
	122010	<u> </u>	1.005		1	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4643	35011	Ā	4684	1	3252	PPGPERSRLGLGVSLHQRSCPK
						CIAVFTRVSEPRIQFPASRILPSS
l						NTSKDFDPVSGQSNYGGSQGS
						GQTLNRPPVASNPVTPSLHSGP
İ						APRMPLPASQNPATTPMPSSSF
						LPEANLPPPLNWQYNYPSTASQ
						TNHCPRASSQPTVSGNTSLTTN
						HQYVSSGYPSLQNSFIKSGPSVP
						PLVNPPLPTTFQPGAPHGPPPAG
						GPPPVRALTPLTSSYRDVPQPLF
		ĺ				NSAVNQEGITSNTNNGSMVVH
	1					SSYDEIEGGG
4644	35012	В	4685	51	236	
4645	35013	Α	4686	1004	1405	
4646	35014	Α	4687	1	771	
4647	35015	Α	4688	1	405	SENVDDVSVMMG/TPANKALL
						DTTGFWHD/DFNNATPNDICVA
						IRSE/AADAGIAQAIMQQLEEA/
						LKQQA/LDRNLNVMMFSDNVT
						LE/DEIQLK/TRAREKGLLV/MG
	}				,	PDCGTSMIAGTPLAFA/NVMPE
				:		GNIGVIGASGTGFRR
4648	35016	В	4689	1	1656	
4649	35017	Α	4690	1657	2259	LPQPQPATPWPSAPTPRFASPAA
ŀ						AMATLWSGTCRIRLWSGSSRA
						TRTAPAALIFPITALGSGQGAW
						TTRCAAGTCGRAASCSSMTSA/
						AQIFSPCHCPNQDWLAVGMESS
						NVEILHVGKPEKYQLHLHESCV
						LSLKFAPCGRWFVSTGKDNLL
						NAWRTPYGASIFQSKESSSVLS
						CDISRNNKYIVTGSGDKKATVY
						EVVY

534

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4650	35018	Α	4691	62	1371	QNQSTISVMLRSAMTTTMAAD
						LITAI/MGQRKQHIDTGFWHDD
						FNNATPNDICVAIRSEAADAGI
						AQAIMQQLEEALKQLAQGSGS
						SQALTQVRRWDSACQKLPDAN
		1				LALISVAGEYAAELANQALDR
						NLNVMMFSDNVTLEDEIQLKT
						RAREKGLLVMGPDCGTSMIAG
		ŀ				TPLAFANVMPEGNIGVIGASGT
				İ		GIQELCSQIALAGEGITHAIGLG
						GRDLSREVGGISALTALEMLSA
						DEKSEVLAFVSKPPAEAVRLKI
ł						VNAMKATGKPTVALFLGYTPA
						VARDEN/VWFGSSLDEA\SLAG
						CFSVRARSPLTRIDGMMILGMF
				1		GGCFAASLWANNVKLRMPRSR
						IRIMQAIIGGIIAGFGARLAMGC
						NLAAFFTGIPQFSLHAWFFANP
		ŀ				LLIGQTLEDPHEYIDYLDKEFPL
		ŀ				YQLVECVVSLNYSYHWECTEI
4651	35019	Α	4692	1	1125	MEAEVDKLELMFQKAESDLDY
		ŀ				IQYRLEYEIKTNHPDSASEKNPV
						TLLKELSVIKSQYQTLYARFKP
						VAVEQKETKSRICAGMTKTMN
		j				VIQKLQKQTDLDLSPLTKEEKT
		İ				AAEQFKSHSFGMWPCKLKYRQ
						NKKKKKKLSQNRSTTWKLNNL
						LLNDYWIQNEMKAEIKMFFET
				ļ		NENKDTTYQNLWDTFKAVCRG
		ł				KFIALNAHKRKQERSKIDTLAL
İ		ł				QLKELEEQEQTHSKASRRQEIT
						KIRAELKEIETQKNLQKINESRS
						CFFEKINKMDRLLARLIKKKRE
			!			KNQIDAIKNDKGGITNDPTEIQT
ŀ						TIREYYKHLYANKLENLEEMD
						KFLDTYNLPRLNQEEVESLNRPI
		$oxed{oxed}$				TGSEIEARINSLPT\KKSPGPDGF
4652	35020	Α	4693	2	421	GRVGGRVGKIRT*LN*IETKKY
						KR*NETKSWFFEKIKMDRPLAR
1						LTKKRIKEIQITSLRNETGNIITD
					1	TTEIHKIIQG*SSSSSSSSSSSSSSS
						SS\FSSSSSSSSSSSSSDTFKRPIT
						GIKIEMVF*KLPTKKSRISL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	-	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
4653	35021	Α	4694	1	654	MEYYAAIKNNTFMSFAATWM
						KLEAIILSKLTKEQKTKRHMFSL
						VSTLELGALTALVRCGRWRRQ
						CDDEAMACVTQGRPLARLTKK
		ŀ				RREKIQITSLRNETGDITTDTTEI
						QKIIQGYDEHTYAHKVENLEG
						MDKLLEKYNPPRLNQEELGTL
						NRPITSYEMEMIIKKL/PNEKSP
		ŀ				GPDGFTAEFYQTFKEELVPILLA
		ŀ				LFHKTEKEGILPNSFYEASITL
4654	35022	A	4695	1	786	MPPSLAYKNPRPQQADTQVA/T
4034	33022	A .	4093	'	780	MSRGVHWRKKMQAAGPREDV
						1
						KRSTQAEEHTDRHQHTSR\HQL
						LLMRIFEMGWKKPSPFQEESIPI
1						ALSGRDILARAKNGTGKSSAHD
1						IPLLKRLDLKKDTIQTIVIVPTG
						GPALQVSQICIQVSKHMGGVK
		ļ				VVMTTGGTNSGDDVLRLDDTV
						HMVIAAPGRILNLIKKGVAKLE
						ETYLRHIGRPGHFGHFGLAINLI
						TYGDHFNLKGIEEQLGKEIKPIP
<u></u>						SNIDKSLHVAEFHSKAVENEKP
4655	35023	В	4696	1	501	
4656	35024	Α	4697	2	573	YSACFFLFSIAMGILLTVPPSFWI
						PTSFSAFLGFFSSFSLLVLHQPD
						FSFVLGLWRIISLLVLKIVEGSS
						NQGMQMASRSWEWPSSSRKM
		1				ATSVRTLPEP/GPSGCRAPSAFPF
1						RKEAGADPSGCPGGRQVPLVAI
		ŀ				GRGGALEPQRWELRAPGSAGR
		ŀ				LPREGGRTFPGAQSPAGAQSPA
						GKQSPPGAQSPLH
4657	35025	Α	4698	2	346	PPINISVPHC*PFGG/EPLEILIPAP
						ERSSHVLSQSPVRTHSSAVHQS
		1				VGASLNCGDKQPPNFSGSKIFF
]				LIYLHLMTGQVRGSSVLCHPNT
						GIQKEGGTVNEIPIAIEKRKKHA
4658	35026	В	4699	1	468	
4659	35027	Α	4700	2	284	ETGEFTQLKELNIQGNCWTLLL
						PELGNLYLTGQKKVCKVENSP
						WVTPIAGQFQLDVSCVSECVCS
						ETYEYLYGQHMQ\ANP\EPPKH
			L_	<u></u>		NNHKSGKD
4660	35028	Α	4701	5	189	
4661	35029	Α	4702	38	190	

NO: of peptide sequence hod in USSN 09/540,217 location of first codon for peptide sequence hod of peptide sequence hod of peptide sequence hod deletion, \=possible nucleo hod sequence hod of peptide sequence hod deletion, \=possible nucleo hod sequence hod of peptide sequence hod deletion, \=possible nucleo hod deletion, \=possible nucleo hod sequence hod of peptide sequence hod hod sequence hod deletion, \=possible nucleo hod deletion, \=possible nucleo hod hod sequence hod hod sequence hod hod sequence hod hod sequence hod hod sequence hod hod sequence hod hod sequence hod hod sequence hod hod sequence hod hod sequence hod hod sequence hod hod sequence hod hod sequence hod hod sequence hod hod hod sequence od hod hod hod hod hod hod hod hod hod	SSFDPGVA ASEVPCASG ETTYRKTT VGSLDPTP GADGSLLH QAEFLQKL LLPKFYGL VMKNLLLR DILQSYRFV GDTVSVHR ITAFKKIPA
4662 35030 A 4703 2 882 WPSVSSG/PSSAVSF SCTSSSASGIQRPMA IPIKKIGHRGVDSSG SSALKCAIQLCITHT ERHVLIEPLIELSSSG VSIHDEFIIKTVQHK IPGYHIDLNQNSWT CCVKAGGKNTQIAV LNSEGERILLCIGITI KKLEHSWKALVCD PGFYTERFQCFMCN RTLTTPNAGKDVEG GMRSGTATVEDSLV 4663 35031 A 4704 2 410	SSFDPGVA ASEVPCASG ETTYRKTT VGSLDPTP GADGSLLH QAEFLQKL LLPKFYGL VMKNLLLR DILQSYRFV GDTVSVHR ITAFKKIPA
4662 35030 A 4703 2 882 WPSVSSG/PSSAVSF SCTSSSASGIQRPMA IPIKKIGHRGVDSSG SSALKCAIQLCITHT ERHVLIEPLIELSSSG VSIHDEFIIKTVQHK IPGYHIDLNQNSWT CCVKAGGKNTQIAV LNSEGERILLCIGITI KKLEHSWKALVCD PGFYTERFQCFMCN RTLTTPNAGKDVEG GMRSGTATVEDSLV 4663 35031 A 4704 2 410	ASEVPCASG ETTYRKTT VGSLDPTP GADGSLLH QAEFLQKL LLPKFYGL VMKNLLLR DILQSYRFV GDTVSVHR ITAFKKIPA
SCTSSSASGIQRPMA IPIKKIGHRGVDSSG SSALKCAIQLCITHT ERHVLIEPLIELSSSC VSIHDEFIIKTVQHK IPGYHIDLNQNSWT CCVKAGGKNTQIAN LNSEGERILLCIGITI KKLEHSWKALVCD PGFYTERFQCFMCN RTLTTPNAGKDVEC GMRSGTATVEDSLN 4663 35031 A 4704 2 410	ASEVPCASG ETTYRKTT VGSLDPTP GADGSLLH QAEFLQKL LLPKFYGL VMKNLLLR DILQSYRFV GDTVSVHR ITAFKKIPA
IPIKKIGHRGVDSSG SSALKCAIQLCITHT ERHVLIEPLIELSSSG VSIHDEFIIKTVQHK IPGYHIDLNQNSWT CCVKAGGKNTQIAN LNSEGERILLCIGITI KKLEHSWKALVCD PGFYTERFQCFMCN RTLTTPNAGKDVEG GMRSGTATVEDSLN 4663 35031 A 4704 2 410	ETTYRKTT VGSLDPTP GADGSLLH QAEFLQKL LLPKFYGL VMKNLLLR DILQSYRFV GDTVSVHR
SSALKCAIQLCITHT ERHVLIEPLIELSSSC VSIHDEFIIKTVQHK IPGYHIDLNQNSWT CCVKAGGKNTQIAN LNSEGERILLCIGITE KKLEHSWKALVCD PGFYTERFQCFMCN RTLTTPNAGKDVEC GMRSGTATVEDSLN 4663 35031 A 4704 2 410	VGSLDPTP GADGSLLH QAEFLQKL LLPKFYGL VMKNLLLR DILQSYRFV GDTVSVHR ITAFKKIPA
ERHVLIEPLIELSSSC VSIHDEFIIKTVQHK IPGYHIDLNQNSWT CCVKAGGKNTQIAN LNSEGERILLCIGITI KKLEHSWKALVCD PGFYTERFQCFMCN RTLTTPNAGKDVEQ GMRSGTATVEDSLN 4663 35031 A 4704 2 410	GADGSLLH QAEFLQKL LLPKFYGL VMKNLLLR DILQSYRFV GDTVSVHR ITAFKKIPA
VSIHDEFIIKTVQHK IPGYHIDLNQNSWT CCVKAGGKNTQIAV LNSEGERILLCIGITI KKLEHSWKALVCD PGFYTERFQCFMCN RTLTTPNAGKDVEC GMRSGTATVEDSLV 4663 35031 A 4704 2 410	QAEFLQKL LLPKFYGL VMKNLLLR DILQSYRFV GDTVSVHR ITAFKKIPA
IPGYHIDLNQNSWT CCVKAGGKNTQIAN LNSEGERILLCIGITI KKLEHSWKALVCD PGFYTERFQCFMCN RTLTTPNAGKDVEC GMRSGTATVEDSLN 4663 35031 A 4704 2 410	LLPKFYGL VMKNLLLR DILQSYRFV GDTVSVHR ITAFKKIPA
CCVKAGGKNTQIAN LNSEGERILLCIGITI KKLEHSWKALVCD PGFYTERFQCFMCN RTLTTPNAGKDVEQ GMRSGTATVEDSLN 4663 35031 A 4704 2 410	VMKNLLLR DILQSYRFV GDTVSVHR ITAFKKIPA
LNSEGERILLCIGITE KKLEHSWKALVCD PGFYTERFQCFMCN RTLTTPNAGKDVEQ GMRSGTATVEDSLV 4663 35031 A 4704 2 410	DILQSYRFV GDTVSVHR ITAFKKIPA
KKLEHSWKALVCD PGFYTERFQCFMCN RTLTTPNAGKDVEQ GMRSGTATVEDSLV 4663 35031 A 4704 2 410	GDTVSVHR ITAFKKIPA
PGFYTERFQCFMCN RTLTTPNAGKDVEQ GMRSGTATVEDSLV 4663 35031 A 4704 2 410	ITAFKKIPA
RTLTTPNAGKDVEC GMRSGTATVEDSLV 4663 35031 A 4704 2 410	
GMRSGTATVEDSLV 4663 35031 A 4704 2 410	OELSSLI M
4663 35031 A 4704 2 410	(<
	/VSYKTKH
4664 35032 A 4705 2 729	
4004 33032 A 4703 2 728	
4665 35033 A 4706 I 1208 MKMEKVNTSWLLP	ľ
RGAGSMVLLLQSQI	RQYIFEYDS
SDRLLAVTMPSVAF	CHSMSTHTS
IGYIRNIYNPPESNA	ľ
DGRILKTSFLGTGRO	QVFYKYGK
LSKLSEIVYDSTAV1	rfgydett
GVLKMVNLQSGGF	
GPLVDKQIYRFSEE	GMVNARFD
YTYHDNSFRIASIKF	'VISETPLPV
DLYRYDEISGKVEH	iF\IITTAEM
TLSKHFDTHGRIKE	
RSLMYWMTVQYDS	
	ľ
QLQSVACNDRPTWI	
	1
DRITRLGDVQY/KID	`
RGSDIFEYNFNGAP	`
GILQSLQGLHEVHR	
4666 35034 A 4707 1 663 MMLLLLALLGAGL	
WHAPARNKIPRAQI	E E
DPKPILELPLAELAC	
LESILCSYLKQALKV	
MDFLGECEEELQAL	
RGLLYGVPMSLKD7	
KPATKDGVIMKVLF	-
KTNIPLTLLRSLKRA	I I
TPIYGQMLSPLNLKI	
GDHGGWPHGPGRG	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4667	35035	A	4708	2	624	 SDARLHKDDTDICFSKTLNSCK
1007	35055	ļ.,				VPQIRYASVERLLERLTDLRFLS
		1				IDFLNTFLHTYRIFTTAAVVLGK
						LSDIYKRPFTSIPV\RRARKLSLT
						SPLNSKIGALDLTTSSSPTTTTQ
		1				SPAASPPPHTGQIPLDLSRGLSS
		İ				PEQSPGTQPEVSGSSPHSTAQSK
		ļ				IWSLVWKQYWLMAPSHALKT
		1				CHAARLARTFVTSSSATKVHCA
		ł				ISLK
4668	35036	A	4709	1	195	ISEK
4669	35037	A	4710	1	1845	MAEAEPGRPGERGGGGAGRAG
4009	33037	^	4/10	1	1043	GRPGGGGGMAEPSGAETRPPIR
						VTVKTPKDKEEIVICDRASVKE
-						FKEEISRRFKAQQDQLVLILSGK
					r	ILKDGDTLNQHGIKDGLTVHLV
					j	IKTPQKAQDPAAATASSPSTPDP
	•		}			ASAPSTTPASPTTPTQPSTSDSA
			ı			SSDAGSGSRRSSGGGPSPGTGE
						GSPSATASILSGFGGILGLGSLG
						LGSANFMELQQQMQRQLMSNP
						EMLLQIMENPLVQDMMSNPDL
	ŀ	1				MRHMIIAKPQMQQLMERNPEIS
						HMLNNPELMRQTMELARNPAV
:						MQEMMRNQDRALSNLESIPGG
						YNALRRMYTDIQEPMFSAARE
		1				QFGNNPFSSLAGNSDSSSSQPLR
		l				TENREPLRNP/WSPSPPTSQAPG
						SGGEGTGGSGTSQP\GSGMFNS
		1				PEMQALLQQISENPQLMQNVIS
		İ				APYMRSMMQTLAQNPDFAAQ
						MMVNVPLFAGNPQLQEQLRLQ
						LPVFLQQMQNPESLSILTNPRA
		1				MQALLQIQQGLQTLQTEAPGL
						VPSLVSFGMSRTPAPSAGSNAG
						STPEAPTSSPATPATSSPTGASST
		-	1			QQQLMQQMIQLLAGSGNSQVQ
						TSEVRFQQQLEQLNSMGFINRE
		<u> </u>				ANLQALIATGGDINAAIERLLGS
4670	35038	C	4711	59	464	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4671	35039	Α	4712	1	1902	MAEAEPGRPGERGGGAGRAG
1				1		GRPGGGGGMAEPSGAETRPPIR
		•				VTVKTPKDKEEIVICDRASVKE
						FKEEISRRFKAQQDQLVLILSGK
						ILKDGDTLNQHGIKDGLTVHLV
						IKTPQKAQDPAAATASSPSTPDP
						ASAPSTTPASPTTPTQPSTSDSA
						SSDAGSGSRRSSGGGPSPGTGE
						GSPSATASILSGFGGILGLGSLG
						LGSANFMELQQQMQRQLMSNP
	İ					EMLLQIMENPLVQDMMSNPDL
		l				MRHMIIAKPQMQQLMERNPEIS
						HMLNNPELMRQTMELARNPAV
						MQEMMRNQDRALSNLESIPGG
						YNALRRMYTDIQEPMFSAARE
						QFGNNPFSSLAGNSDSSSSQPLR
						TENREPLPNPWSPSPPTSQAPGS
	1					GGEGTGGSGTSQVHPTVLNPFG
						INAASLRSGMFNSPEMQALLQQ
						ISENPQLMQNVISAPYMRSMM
		l			•	QTLAQNPDFAAQMMVNVPLFA
		1				G\NPQLQEQLRLQLPVFLQQMQ
ľ		l				NPESLSILTNPRAMQALLQIQQ
		}				GLQTLQTEAPGLVPSLVSFGMS
						RTPAPSAGSNAGVYPPRPPLPHP
						ATPSHIFSNRG/SPAPQQQLMQQ
		1		,		MI\QLLAGSGNSQVQTPEVRFQ
						Q\QLEQLNSMGFINREANLQALI
						ATGGDINAAIERLLGSQLS
4672	35040	В	4713	309	527	

539

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4673	35041	A	4714		2506	KKDQRETRNQKGLGFLRDRPQ WVSGKMKQECLQ/HWNAMAI NNHRAVAIFPKRRHGREDGLLS NLPTFGSYAPLRRVSNEFIASAV QRILLERRALPHGNCKLGGLTP QFMNRKNALVLEVLARAIRQE KEIKGIQLGKEEVKLSLFADDM IVYLENPIVSAQNILLKLISNFSK VSGYKINVQKSQAFLYTNNRQ TESQIMSELPFTIASKRIKYLGIQ LTRDVKDLFKENYKPLLNEIKE DTNKWKNIPCSWVGRINIVKM AILPKVIYRFNAIPIKLPMTFFTE LEKTTLKFIWNQKRARIAKSILS QKNKAGGITLPDFKLHYKATV TKTAWYWFQNTDIDQWNRTEP SEIMPHIYNYLIFDNPEKNKQW GTDFLFNKWCWENWLAICRKL KLDPFLTPYTKIKSRWIKDINVR PKTIKALEENLGNTIQDIGMGK DFMSETPKAMTTKVKIDKWDL IKLKSF
4674	35042	A	4715	3	372	SVGVALRPWKRE/RAASERRSS SGGGGGGGGGGGGGGGSGS GQ/HAPAAPAGGIEAVNMASAS YHISNLLGVLAAAALAVTQALP \PPSAVAGSFSAPKSPAHRSAGL PIPAEPLSSPLLQPPPP
4675	35043	Α	4716	1	1008	11112123012221111
4676	35044	A	4717	1	2619	
4677	35045	Α	4718	449	801	VLLLSSMSRRKCQSLYVDLLM KKETE*SMEKEKLTMHPLSCTH I*PRPQAPEADKQMRR*TEEQN CRMMWQKKEKEHLNAKRSLA GSGWRDQPLDGKAPGEDHLPIP SPFQLPIHPI
4678	35046	Α	4719	1	1255	
4679	35047	A	4720	2	843	CLHGFGYRIRDSELQKIHRAAV KGDAAGVERCLARRSGDLDAL NK/TAQIAGAQPREEACTVILLE HGTNPNLKDIYRNTALHYAVY SESTSLAEKLHFHGANIEALDK VLSISFLSKILMSSLKTCGRDAE DYTISHHLTKIQQQILERKKKIL KKEKRGKASESEFLNSLGGPTL DKKIRNVEISDESAVSILHELCV DSLPALDDEVLSVATKCVPEKV SEPLCRPSHEKGNRIVNGKGEG SEECLCPAAHRLRCGERLYLPP RLGCERLCLATTPSEK
4680	35048	Α	4721	295	1050	

SEQ ID NO:	SEQ ID NO: of peptide sequence	t	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4681	35049	В	4722	306	554	
4682	35050	A	4723	2	7973	
4683	35051	Λ	4724	2	316	
4684	35052	Α	4725	81	228	DPPLCLGLL\LHRASIVQKRIIYF
						QDEGSLTKKLCEQGKTLKHSQ QMFFK
4685	35053	Α	4726	1	1043	MELEWKVNVKVNSQDTNHHG
	į	1				SLQLARGEAAAVKFGRMSKKQ
						RDSLYAEVQKHQQRLQEQRQQ
		l				QSGEAEALARVYSSSISNGLSN
						LNNETSGTYANGHVIDLPKSEG
						YYNVDSGQPSPDQSGLDMT\GI
						KQIKQEPIYDLTSVPNLFTY\SSF
		ŀ				NN\GQLAPGIT\MTEIEIVAAEKF
		1	1			PLIYKQSFLLTVLSFGGGGSVIC
		l				GPTFAKVSSSRRFIHHGDKIQPS
						INALGWTFMEETPQIFKCRTNT
		1	İ			HGKELEHDLPEHSSGQGSTRRK
						SSCLRRDNNPMLLSGGRFYEKI
						HNFITGTFDVRKMEHAEGKTSD
						LVHVGFQAIKMPSSLKQEASNG
			i			LIKLEEASGARMKTGHIK
4686	35054	Α	4727	467	584	
4687	35055	Α	4728	1	1794	
4688	35056	Α	4729	110	1797	PSQQEPGSGTSCLRYCWTAQTL
ŀ		1				PSVTMKLWVSALLMAWFGVLS
						CVQAEFFTSIGHMTDLIYAEKE
l						LVQSLKEYILVEEAKLSKIKSW
						ANKMEALTSKSAADAEGYLAH
						PVNAYKLVKRLNTDWPALEDL
						VLQDSAAGFIANLSVQRQFFPT
					:	DEDEIGAAKALMRLQDTYRLD
			,			PGTISRGELPGTKYQAMLSVDD
ŀ		1				CFGMGRSAYNEGDYYHTVLW
						MEQVLKQLDAGEEATTTKSQV
<u> </u>						LDYLSYAVFQLGDLHRALELTR
						R\LLSLDPSHERAGGN/LCRYFE
						QLLEEEREKT\LTNQTEA\ELTTP
						EGIYERPVDYLPERDVYESLCR
					,	GEGVKLTPRRQKRLFCRYHHG
						NRAPQLLIAPFKEEDEWDSPHI
				1		VRYYDVMSDEEIERIKEIAKPK
						LARATVRDPKTGVLTVASYRV
						SKSSWLEEDDDPVVARVNRRM
						QHITGLTVKTAELLQVANYGV
						GGQYEPHFDFSRNDERDTFKHL
						GTGNRVATFLNYMSDVEAGGA
						TVFPDLGAAIWPKKGTAVFWY
						NLLRSGEGDYRTRHAACPVLV GCKWVSNKWFHERGQEFLRPC
	ī					

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
4690	35058	A	4731	425	1079	PMGTSFLYECANFQSVSLGDPQ
	}	1				EPSCLCGGQPQGAAGGSSGSED
						ALLCLWDLQARKWKFEMSYTS
		1				GNIRGSSFCQFKSSFLVYFKALL
						KFRELFAQFKSFMGEKKLSLAP
						PMSDSSGDGIIRKIDYASCKRSG
						\SSIRALPKTYQQP/EVLNDTWV
						SFPSWSEDSTFVSSKKTPYEEQL
						HRCEDERFEEMKTAGFASFGER
						SKTGVPNLRVTDWYRLVAC
4691	35059	Α	4732	3	425	GASSEEAEGASEGP\GAPGGWG
						APGSQGAQEGGDLQEAEESQE
						GGDPRKPRSPRKVERHRKAGA
		1				PGRDLGRPSLTVLL\NHCVLQR
						LRKIYHSSIKPLEQSYKYNELRQ
						HEITGQRCPCCEEPKPQHQERA
						LFVVVSENRNI
4692	35060	В	4733	1	1056	
4693	35061	Α	4734	171	511	LLSVRHVVRNTQETANDVQVW
			•			/LDREGGSKI\NTGVCFLDHMLD
	1			-		QIATAVSRME\INVKGDLYIDD
						HHTV\EDTGL\ALGEALKIAPGD
	1					KPGICRFGFVLP\MDECL\ACAL
	<u>.</u>	<u> </u>				DISGGPH
4694	35062	Α	4735	563	763	
4695	35063	Α	4736	365	1644	RTSQMSSSAWRQQNRARPSSAI
						LPSSLSLGHPALPQFSQRMPAT
						ASQLPGMVGVLEGYGQTAASP
						GSVSSCSPACSSCCLGCWWPSS
	ĺ					WPSCRLHPHLGRPIAHCLPE/VL
						TTTTTTTTTTTSQAAGTPKGQQE
						SGVSPSPQSTCGGLLSGPRGFFS
						SPNYPDPYPPNTHCVWHIQVAT
						DHAIQLKIEALSIESVASCLFDR
						LELSPEPEGPLLRVCGRVPPPTL
	:	İ				NTNASHLLVVFVSDSSVEGFGF
						HAWYQAMAPGRGSCAHDEFR
			1			CDQLICLLPDSVCDGFANCADG
						SDETNCSAKFSGCGGNLTGLQG
		1				TFSTPSYLQQYPHQLLCTWHIS
						VPAGHSIELQFHNFSLEAQDEC
						KFDYVEVYETSSSGAFSLLGRF
						CGAEPPPHLVSSHHELAVLFRT
						DHGISSGGFSATYLAFNATERL
4606	25064	-	4727	1	154	CLVESTSS
4696	35064	Α	4737	11	154	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			}	sequence		
		<u> </u>				
4697	35065	Α	4738]1	700	GSRPQFPGHTRVRASGWRPCSL
]				l		KPQLLGPVHPVSPYLLFLSSSDC
						AMGLIDGEWQLVLNVWGKVE
i						ADIPGHGQEVLIRLFKGHPETLE
						KFDKFKHLKSEDEMKASEDLK
				ļ		KHGATVLTA\LGGILKKKGHHE
	1					AEI\KPRGTVSNATKHKIPVKVT
						WRSSPECII\QVLQSKHPGDF\GA
						\DAQGAMKQGPGSCFRKDNWP
						PTYKELGLSRAKPLAGFPTPNPS
						WAPGFKRERGLISV
4698	35066	Α	4739	1	154	
4699	35067	Α	4740	1	617	GSRPQFPGHTRVRASGWRPCSL
						KPQLLGPVHPVSPYLLFLSSSDC
					,	AMGLIDGEWQLVLNVWGKVE
ŀ	1			1		ADIPGHGQEVLIRLFKGHPETLE
						KFDKFKHLKSEDEMKASEDLK
ļ	1					KHGATVLTA\LGGILKKKGHHE
1						AEI\KPRGTVSNATKHKIPVKVT
		1				WRSSPECI\IQVLQSKHPGDF\GA
						\DAQGAM\NKALE\LFRKD\MAS
						NYKELGFQG
4700	35068	С	4741	46	522	
4701	35069	Α	4742	78	617	TKELLHSKRNCHQSEQATYKM
				1		GENFCNLLI*QSANIQNLQRT*T
		1		İ		NLQEKNKQPHQKVGEGYEQTL
				'		LNRRSLCSQKTHEKILIITGHQR
ł						NANQNHNEIPSHTS*NGNH*KV
						RKQQVSYKLL*MRPRRTRQVT
						Q*RREPETSLAKETPGNPTNTN
						AKFKTRGARISHYSSGNGERLP
1		1				RTVC
4702	35070	В	4743	1	6477	
4703	35071	Α	4744	1	623	MSSDISEVEDKNEFLTEQLSKP
1		1				QIKFNTLKDKFLKTRDTL\RKKS
1		1	1			LALETVHNNLSQTQQQIKEMK
		1				EMYENAEAKENNSTGKWSCVE
1		1	ļ			ERICQLQHENPCIEQQLDDVHQ
1		1				KECLPSRKEKFKSEPPAFLSGN
						QVKSSSCSLQTLFPPDDLILYLE
						NPKDSTKKLLELINKFRVTGYK
	1					IKLQKSVAFLNDKNEQSKEENQ
						ECNPIYNNYK
4704	35072	A	4745	2	3272	
4705	35073	A	4746	1	579	
4706	35074	A	4747	3	510	
1700	1330,4	l' .	L'''' —	l	12.0	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
4707	35075	Α	4748	li i	1261	MEAVDTFLVFNALNFLLTSQTT
	:					GCSSSCDPGHGLGQGSGNRFPI
				•		AAASARTAEEAKWFFSNKDYS
						NVLQLPDPKKTEYHIHEKRFSD
						SRILCYYPEFGKVEEIILTAMKH
						DWFGKHRKDDKIEKTGKIKIQE
						SFTSEEERIRMKQEQERIQAKTR
		1				EFRE\RQARERDYAEIQDFHRTF
						GCDDELMYGGVSSYEGSMALN
						ARPQSPREGHMMDA\LYAQVK
						KPRNSKPSPVDSKGKVLGEADP
						VYPRRNPGTEGNASSLPSVSFSE
						SPPVRAAFAGYHYVAFPWLQK
						LMSGTSGKDPEKEADLARVVT
						LRVVTLRIETHVKTLAFTAALL
						QQLACIDVGMDKQNDVYTYN
ŀ	,	l				KILLSFKRKEILTHTTTRMSLED
						IILSEINQWQEDKYRFGLYEVA
						QVVKLIETESKVVVSRDGGRG
4708	35076	Α	4749	10	2051	Q V V NBIE 125K V V BIND GENO
4709	35077	A	4750	2	2118	
4710	35078	Α	4751	1	658	MWNSKTLAAFRPCPKDPLNFE
						LERDNLAYLAEEIPKQQSIQYIT
ļ						WMILKAFSHMHLQRDNLKLEL
						MFKRKAKHKGLKNLHPDHVIE
						KKNLFSAEKFKPAAEIYISNEEP
						NVNSQDNGKKCLQGMSEIFAA
						APAITONTSDKTTLIKVSSWPYI
						ADRRCPLVNVTREDSPSEDPVF
						LRTLGKGDWFGEKALQGWGL\
			8			KGIHHVSAQEPVCLLLPFMTPR
4711	35079	В	4752	1	471	
4712	35080	Α	4753	315	407	
4713	35081	A	4754	411	1042	
4714	35082	Α	4755	1 .	423	
4715	35083	С	4756	202	321	
4716	35084	Α	4757	5	413	CFFFFFFETESHFVTQAGVQWR
						DLGSLQSPPPGFTPFS/SPQPPKE
1						PGPQAPATTPGQSFAFLVEMGF
						HHVSQ\EVSIS*PRDPPASASQS
						AGTTGVSHRAWTFCFLRQSLA
						LSPDWSAVARSQLTATSASWV
						QVSRR
4717	35085	С	4758	150	491	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4718	35086	I _A	4759	11	548	GIAFCLNLIKTLKLPNFKSCVIL
						LGLLLLYDV/FFVFITPFITKNGE
						SIMVELAAGPFGNNEKNDGNL
						VEATGOPSAPHEKLPVVIRVPK
	ŀ					LIYFSVMSVCLMPVSILGFGDII
						VPGLLIAYCRRFDVQTGSSYIY
	į					YVSSTVAYAIGMILTFVVLVLM
						KKGQPALLYLVPCTLITASVVA
4719	35087	В	4760	642	1985	
4720	35088	Α	4761	39	252	
4721	35089	Α	4762	1	783	
4722	35090	С	4763	218	358	
4723	35091	В	4764	372	374	
4724	35092	В	4765	129	1036	000011105555
4725	35093	Α	4766	1211	1983	SQSCLLLQEDFAPIAGEQEAEQ
						HQEDLRALLRASLQGQCSRQP
						GTRLHGSAPWPGEAQNRSRPLP
						GDSPSLDRYRGI/SDAVGKSRSG
						DIGSSLRVEAGDKRTQASPERQ
						PHCGAHDAQDISGGREIFKPRQ
						LPGSAIWSIKVGHGSGFPGKRR
						PRGAGLSGRGGRGRSKLKSGIG AVVLPGVSTADISSNKDDEENS
						VLDMVVLFSSSDKFTLN/QVCG
						SFGQGAEGRLLACSQCGQCYH
		İ				PYCVSIKMDACSSSELKY
4726	35094	A	4767	1	603	MANFNDCVLDKEKVCIAAKFIT
1,20	3307.		""			HAPAGEFNEVFSDIRLLCNNDS
						LLRERAARAFAHYNMDQFTPV
	Ì					KMEGCEDQTIIACIESHECQPKN
						FWNGRWRSEWKFTITPPTAQV
						VGVLKIQVHYYEDGSVQLVSH
·			}			KDVQDSLTVSNDAQTAKEFIKII
			}			ENAENEYQTAISENCQTMSDTT
						FKVLRRQLPVTRTKIDWNKI\LS
						YNI
4727	35095	Α	4768	1	867	MADFDDRVSDEEKVRIAAKFIT
-		ĺ				HAPPGEFNEVFNDVRLLLNND
		1				NLLREGAAHAFAQYNMDQFH
						AVK\IEGYEDQVLITEHG\DLGN
						SRVLDPRNKISF\KFDHLRKEAS
						DPQPEEADGGLKSWRESCDSA
						LRAYVKDHYSNGFCTVYAKTI
						DGQQTIIACIESHQ\FQPKNFWN
						GR\WRSEWKVP\ITPPSAQ\VVG
						VLKIQVHYYEDGNVQLVSHKD
						VQDSLTVSNEAQT\AKEFI\KIIE
						NAENEYQTAICGNYQTMSDTT
						FKALRRQLPVTRTKIDWNKILS
		<u></u>	<u></u>			YKIGKEMQNA

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	,			sequence		
4728	35096	Α	4769	2	690	WPSGSASRPRHLRAPEPV\GKA
						GKNKGAGLTPA/AGTSGCAGR
				į		AESRGPPTGSHGSEPSRGSKGL
						CGRGTRSLPERGAEVLPHLPAQ
						TSPQLGSQGAWP/RHPRKPFRP
						AARAGTQPPGFVPSTPALLKVL
						YRSAHTQHSGRAAPEPSRLALG
	1					PTLQQKRHMGHRRNLTFSSTY
						NKKCGFGWAVEMRFSALLDVT
						TTPSRVAFWGSRGPPLGMERQL
						QLLSGYSPDGWLSTSM
4729	35097	Α	4770	1	590	MDTRIGTTDTGTYGKVEGARR
		İ				MRLKKLPIEYYAYYLDDEIICTS
						NPCEFPGYGIEREYPCETSGPLE
						LTVQEDQGEEPQREELTRKKTS
						KVCRGSPLAWATGVKPCLKKK
						RRRRKEERRKKKEEGEGEGEE
				4		EEEEEEEEEEEEE
						EE/EEEEEEEEEEEEEEEE
						EEEEEESDPCTSWNTSQP
4730	35098	Α	4771	1	288	
4731	35099	Α	4772		237	MRSSVIGPRSHIPSQRKSGIQEE
ļ						EEEEEQEEEEEEEEEEE
						EEEEEEEKGLLDQEAARHLV
	25100	ļ.,	1772	ļ 	704	LPAATQCKPKM
4732	35100	Α	4773	1	794	MASHSSPMGSQYYYGFPGPDS
ł	:					AMHALNTVVSEKDLTLDLSGL
	,					VARNKRCGPHSYSLNTHLLHA CLRLPTQRENTTLKTFIPQGWEI
ŀ						HTDQVEREAECQPGRLKICVHD
						TAQELPLASTARNALLGRNLCP
				İ		FRQSSTTQMPDEIPISLDDRMRP
						PSLKKKK/VVGEEEEEEE\EKEK
				†		EEEEEE/DEGEEEEEEEEEE
						EEEEEEEEGGGGVGEEEE
		İ				EEGEGEGGGGEEDEEE*EEEEE
						EQKKKEKKKKEQEEGGGEGG
4733	35101	Α	4774	115	341	222
4734	35102	A	4775	1	651	
4735	35103	Ā	4776	189	618	SLCHKEAEGGHGKAHVEGKRA
		-				PSNLQPSAPAELSGNNSISF\HHE
					1	EEEEEEEEEEEEEEEE
					1	EEEEEEEEEEEEEDLLYNFI
						YIIFSKMPTKVPSLWDSKLGAE
						QAAPEENNKKEQQEEYQGKSF
					1	SFLNLTECWP
4736	35104	Α	4777	1	414	

SEQ ID			SEQ ID NO:	ľ		Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		07/340,217	sequence	or peptide sequence	deterior, 1-possible nucleonide insertiony
			•	·		
4737	35105	Α	4778	189	692	SLCHKEAEGGHGKAHVEGKRA
į			ļ	1		PSNLQPSAPAELSGNNSISF\HHE
						EEEEEEEEEEEEEEEEE
						EEEEEEEEEEEEEDLLYNF
						IYIIFSKMPTKAMSISSMFKFAT
						KIRMTVTPNPKCCWIPVFPAYL
						STSKAAKCMHHTAHHGQEKSD
						HCISPSPIPAP
4738	35106	Α	4779	2	3815	
4739	35107	A	4780	957	1493	KNRNYNKLSLRPQCNPTRTQD*
						ETHSKPLNYMETEQPSPE*LLG
						T*RNEGRHKAVL*NQ*EKRHNI
						PESLGHI/AKQCVEGNL*H*MPT
		1				RESRKDLKLTP*/PSQLKELEKQ
		1				EQTHSKAGRRQE/TN*DQSRIEG
						DRDTKNPSKNQ*IQELVF*KDQ
						QN**TASKTNKEEKREESNRRN
						KKC
4740	35108	Α	4781	1	4962	
4741	35109	Α	4782	2421	3011	NQACQPGQAGAARAGQCFPRV
						AQRPGPGPAGMALAHPDLYLC
						HSAGDQCR*DEEADPPEGQAPT
1						HFQVHPAGGLHLLDRLAAPQH
1						QAGTQGQGQPGRHH*PAARSS
						PVEPPSTGSPSRPTTTMWPSCSR
						RQGGLGPGL*CVQQPHLLDAP
	1					GTLPVLGALVCRLQPWRCLPL
						GGGGGVGGSQRPPGGGGGNG
						GRNWG
4742	35110	Α	4783	1	932	MRTPKSSIKPSLGEEKENYRGS
						LKETAPSLQKEWAEQSSKSQET
						VGVKGITRHFQVSTLLERREDR
						KSKEPPLLIPRQTVSGVDLQQTP
						TDLQLSVLTVRRKTNKQKGHP
						HQKPICTSPSSNTKVKNLEKQL
						DEWLTRITNAEKSSKDRMELKT
						KARELHDECTSLSSRCDQLEER
			į			VSVTEDEMNEMKRGEKFREKR
						IKRNEQSLQEIWDYVKRPNLCL
					[IGVPESDGENGTKLENTLQDIIQ
						ENFPNLARQANIQIQEI/RENAT
						KILLEKSNSKTHNCQIHQS*NEG
		1				KNVKGSQRERSGYPQREAHQT
4743	35111	Α	4784	477	638	LLCCGFELPLLARRSLIV*SLLLS
7,73	33111	l'`	7,704	' '		THQSHSPSSFVPLLVRSCIPLEEE
						RRSDF
4744	35112	В	4785	1	1509	
7/77	33112	سا	17703	l <u>.</u>	1.507	<u> </u>

SEQ ID			SEQ ID NO:	1	li .	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		05/340,217	sequence	or peptiae sequence	deterion, (-possible nucleonide insertion)
				l		
4745	35113	A	4786	3	832	ERHHPPPQAGPHPPRRRKERNT
						HRQPTPQPHGA/WKISVPQTTN
1]					KGISSSITSTQRNQPQATTVEKA
						HNESTRTPEEPKGAVRGTR*KK
						KKTSRDTPPQTP*EKSSSATRNK
						AGGKDSDKRERKATEDQTTPS
	ŀ		<u> </u>			KRRKLHPMAKKLKTLKKN*MN
		İ				G*LE*PMQRSP*RT*WS*KPRHE
						NYVMNAQASVTDAINWKKGPP
						PRPTTNDAKIVTRPTHPTCSHH
					<u> </u>	RPPRKPPRTHPTPTPTQNKISQ*
						NGYTPPRGKRVREDRCKPPQAP
						TSAPRAAKQRQS
4746	35114	В	4787	1580	4673	
4747	35115	Α	4788	1	462	MKLEHQEAQRRSGQGREKRR
						WPEGKAGPGCEGAWILCPESQ
						DDSKQEGDNNMIVVSRNAVRS
						VKAEFQGDNLENWESCSEAIIV
						QGRDNGVPHKAHGLGTEEEGT
						VLKISERQNWLDLVDFILLGGD
		Ì				LFHENKPSRKTLHTCLELLRKY
						CMGDRPVQFEILSDQSVNFGFR
						KLTNRKDIHTKNPSVRHHHQRP
						KVDETIKMGKTQSRKTRNSKN
						QSTSPPPKERSSSPAIEQSWMEN
						DFDELKEEGFRRSNYSELKEEV
						RTNGKEVKNLEKKLDKWITRIT
						NAEKSLKDLMELKTIAQELRDE
						CTSLSNQCDQLEERVSVMEDQ
						MNEMKREEKFREKRIKRNEQS
						LQEIWDYVKRPNLCLIGVPESD
						GENGTKLENTLQDIIQVNFPNL
						ARQANIQIQEIQRMPQRYSLRR
						ETPRHIIVRFTKVEMKEKMLRT
						AREKGLECSGAGLAHCKLWLL
						GPSDPPDCSSVSPVLRVHLVLPS
		}				SLPHSVGTPFLGSVSIPPSVPRFP
						DRVFHPYPYTHYCDNLKTCHT
						SHGSVMAETAVINHKKRKNSP
						RIVQSNDLTEAAYSLSRDQKRM
						LYLFVDQIRKSDGTLQEHDGIC
						EIHVAKYAEIFGLTSAEASKDIR
						QALKSFAGKEVVFYRPEEDAG
				1		DEKGYESFPWFIKRAHSPSRGL
					_	YSVHINPYLIPFFIGLQNRFTQFR
L	<u> </u>	1	<u> </u>	<u></u>		LSETKEITNPYAMRLYESLCQY

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4748	35116	A	4789	1314	2221	KNRNYNKLSLRPQCNPTRTQD* ETHSKPLNYMETEQPSPE*LLG T*RNEGRHKAVL*NQ*EKRHNI PESLGHI/AKQCVEGNL*H*MPT RESRKDLKLTP*/PSQLKELEKQ EQTHSKAGRRQE/TN*DQSRIEG DRDTKNPSKNQ*IQELVF*KDQ QN**TASKT/IQRRKERRIK*TQ* KMLKEDITTDPTEIKTIIREYYK HLYAHNLENLEEMDKFLDTYT LPRLNQEEAESLNRPITNSEIET VINSLKEKAQDQKDLQLNSTRA LFTIAKAWNQPKCPSMTDEIK/I NVEHIHHGILCSHQKE
4749	35117	A	4790	2	2260	TKDKNHMIISIDGGKAFDKIQQ PFMLKTLNKLGIDGTYLKRIRAI FDKPTANIILNGQKLEAFPLKTG TRQGCPLSPLLFNIVLEVLARVI RQEKEIKGIQFGKEEVKLSLFA DDMTVYLENPIFSAQNLLKLIS NFSNVSGYKINVQKSQAFLYTN NSQIMSELPFTIATKRITYLGIQL ARDVKDLFKENYKPLLNEIKED TNKWKNIPCSQIGRILWPYCPR QEDENFNSLLQNGDILNSSTEE KFKAHDKKDFNLPEYDLNVEE RLVLIEKSVDSTATADDTHKLD HINMNLNKLITNDTFQPEIMERS KTQDIVLGTSFLSINSKEETEHL ENGNKYPNLESVNKVNGHSEE TSQSPNRTEPHDSDCSVDLGISK STEDLSPQKSGPVGSVVKSHSIT NMEIGGLKIYDILSDNGPQQPST TVKITSAVDGKNIVRSKSATLL YDQPLQVFTGSSSSSDLISGTKA IFKFDSNHNPEEPNIIRGPTSGPQ SAPQI\YGPPQYNIQYSSSAAVK DTLWHSKQNPQIDHASFPPQLL PRSESTENQSYAKHSANMNFSN HNNVRANTAYHLHQRLGPARH GEMWAISPNDRLIPAVTRSTIQR QSSVSSTASVNLGDPGSTRRAQ IPEGDYLSYREFHSAGRTPPMM PGSQRPLSARTYSIDGPNASRPQ SARPSINEIPERTMSVSDFNYSR TSPSKRPNARVGSEHSLLDPPG
4750	35118	A	4791	1516	1729	ILAPHSLLACRVSAERSAVSPM GFPLWVTQPFSLAALN/DFFLHF NFG/RI*QLCVLELLFSRSIVVAF SEFP

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4751	35119	A	4792	3	1426	RRYDELQNSSGRDGKPRAMAV TRSTSSTSSGSNSNVLVPVSWK RPQYSQKRAKEKLVHVLSLCG QEVGLSKNPSVIFSSCGDLDLLE HQTSLVSSEDGAREQENMDDT NSEQQFRVFRDFDFLDVELEDG EELQGESMDNFNWGVRRRSLD SLDKCDMQILEERQLSGSTPSL NKMHHEDFDESSEEEDLTASQI LEHSDLIMTLSPSEETNPMELLT TACDSTPPEPHSFNTRMSSFDAS LPDMNNLQISEGSKA\EAVREE EDTTVHEDDLSSSINELPAAFEC SDSFSLDMTEGEEKGNRALDQF TLASFGEGDRGVSPPPSPFFSAI LAAFQPAACDDAEEAWRSHIN QLMCDSDGSCAVYTFHVFSSLF KNIQKRFCFLTCDAASYLGDNL RGIGSKFVSSSQMLTSCS\LDKL KFSVLELQEYLDTYNNRKEATL SWLANCKATFAGGSRDGVITC QPGDSEEKVIKAC
4752	35120	С	4793	60	164	Q1 GDGEDIT VIII 10
4753	35121	В	4794	44	2547	
4754	35122	A	4795	401	9546	PRADITTCDRRITPGTCSRPLPV LPASLYAADMASQQDSGFFEISI KYLLKSWSNTSPVGNGYIKPPV PPASGTHREKGPPTMLPINVDP DSKPGEYVLKSLFVNFTTQAER KIRIIMAEPLEKPLTKSLQRGED PQFDQVISSMSSLSEYCLPSILRT LFDWYKRQNGIEDESHEYRPRT SNKSKSDEQQRDYLMERRDLAI DFIFSLVLIEVLKQIPLHPVIDSLI HDVINLAFKHFKYKEGYLGPNT GNMHI
4755	35123	A	4796	2	6107	GERKPKIDLFRTCVAAIPRLLPD GMSKLELIDLLARLSIHMDDEL RHIAQNSLQGLLVDFSDWRED VLFGFTNFLLREVNDMHHTLL DSSLKLLLQLLTQWKLVIQTQG KVYEQANKIRNSEARDLELDV GVDVDVGVDVDADVDVDVDL GVDVDVGVDVDADVDVDVD MGVDVDVDRIYGFGCKCGCDR DVDRVVDVDVDIDVDVDVDM DVNIDVDVGLGAGVDVDMDV HIDMDMDVDIGSDNILQSALRR

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence	1	02/2/40,217	sequence	or peparae sequence	possible national matrix
		}				
4756	35124	Α	4797	3	565	MLPGHLFCLAKAGKEEQEEKG
			}			QIITRKTITPQNIPGPGAHITATH
						RGRCTVATKFSGGLCKEPNRLT
						WY/GPPLKGAHGGR/GPSSRVPS
						FPAAGQPPIGGALHGAGSRGLA
						RTRAQLGGPPGGELRGT/SGNG
į						DGQGPDGDSRKSGSPLRPSHL/P
						DPAAVAGAPVSGHLQTPGHHA
						GRRQQAGPRGAGLP
4757	35125	Α	4798	1	417	
4758	35126	A	4799	1	1069	MPRPPVPANALGARGEAVRLQ
		1				LQGEELRLQE\ESVRLHQINIYL
						SDRISLHRRLPVRWNPLCKEKK
						YDYDNLPRTS\VTKAFYNEAWS
						TLLRTVYSVLETSPDILLEEVIL
						VDDYSDREHLKERLANELSGLP
						KVRLIRANKREG\LVRARLLGA
						SAARGDVLTFLDCHCECHEGW
						LEPLLQRIHEEESAVVCPVTDVI
İ						DWNTFEYLGNSGEPQIGGFDW
1						RLVF\RWHTVPERERIRMQSPV
						DVIRSPTMAGGLFAVSKKYFEY
						LGSYDTGMEVWGGENLEFSFRI
						WQCGGVLETHPCSHVGHVFPK
						QAPYSRNKALANSVRAAEVW
1						MDEFKELYYHRNPRARLGLAC
						DECSIKAGWWL
4759	35127	Α	4800	1	1152	MHNSDGIEVMRQQKAEIGRVG
						GLEGIQLGVTEVIVNGARMLES
						YNCKAELGATGLVNYQISVKC
						SNQFKLEVYLLNAENKVVDNQ
						AGTQGQLKVLGTNLWWPYLM
						HEHPAYLYSWEGRPDGAQAVG
1						ALTPGTLAV\EVWLTAQKSLGP/
						SDFYTLPVGLRTVAVTESQFLIS
1						GKPFYFHGVNKHEDADIQGKG
						FNWPLLVKDFNLLCWLGANTF
		1		1		CTSHYPYTEEMLQICYRYGIVVI
						DECPAPSGHT\GPSVPSLLARW
i						QLFNNVSMHHHMWVVEEPVL
						RDKNHPAMVMWSLAKEPASFL
				ĺ		ESAGYSFKSLTMEQTARVLDLD
1						TGEAVLQYRSLPRGAHKTLGK
						KRKISSYNVDLTSCQLAKEKCL
						KGPSSFLQSRQERMNSELRDN
4760	35128	Α	4801	293	535	
4761	35129	Α	4802	94	686	
4762	35130	В	4803	1	2187	
	_					

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	j .	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
4763	35131	A	4804	596	1789	MFHVIERPYECKECGKNFRSGY
						QLTLHQRFHT\G!HKGGKPYEC
				·		KECKKTFTLYRNLTRHQNIHTG
						EKLFECKQCGKTYTTGSKLFQH
						QKTHTGEKPYECKECGKAFSL
						YGYLKQHQKIHTGMKHFECKE
						CKKTFTLYRNLTRHQNIHTGKK
						LFECQECGKAYSTGSNLIQHRK
						THTGEKPYKCKECGKTFSLHG
						YLNQHQKIHTGVKPYECKKIHT
						GGKPYECKECGKAFSRASNLV
						QHERIHTGEKPYVCKQCGKTFR
						YGSALKAHQRIHRSIKVNQWL
						DSPRHKSINCDPMSPRTRKTRE
						RSHVSGRSLCQRLLGQGAQLRE
						FKQPPRNRFRRVCGLVARNTFP
		ļ				RGLRGRRASRRNAAASRLEDA
j		İ				VAELLKARCSALGGSGLPEGLS
4764	35132	Α	4805	2	489	CQGEGAKWEQGMTGRRWVVG
1						PPASSTPPHS/HFWLLVLSRGLV
						GIGEASYSTIAPTIIGDLFTKNTR
						TLMLSVFYFAIPLGSGLGYITGS
				1		SVKQAAGDWHCALRESFLQLV
				}		RLLKVTHLLQQLQLLYPTNGK
						RQDTNHTRSQCAHYLPHLDFQ
						VALCAYAVLQC
4765	35133	Α	4806	1	327	
4766	35134	Α	4807	899	1219	ANRKASTMRWYVRPFCSGLST
						LSREKLYPLRMLRMRASTGPLS
						GALGMYSTASPRLMFSGCSISS
						AR*LFRSSAVTMPPAAWMSFTI
<u> </u>		ļ				WAVRLPL*KAWAPSMASVS
4767	35135	Α	4808	2152	2633	SRLDCGQGLVNGSCDYHHGCT
						GDTPRRPEWHRSLAHCMSPVD
ŀ			ł			SLPGCAVSGSSDPWEVSWLWQ
						Q/VDRCDNREE/VMW*QRLGFP
ŀ						SPTVSFPQRAVSASKPLQ*APS
			1			WISLHSTLSQCRASDLRGRHTV
				1		PEVVYEWSPCALPTHREHPNAI
		<u> </u>		<u> </u>		MEGWPPPQAWWA
4768	35136	Α	4809	2	387	SNASVILEGEDLRFSCSVRMAG
:		1				RLQGRFSVIWQLVDRQNRRSDI
		1		-		MWLDR\DGTVQPGSSYWERSSF
		1				GSVQMEQVQPNSFSLGIFNSRK
						EDEGQYECHVTEWVRVVDGE
		1				WQIVGERRASTPISIIALGE

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4769	35137	A	4810	46	3753	EIRSWEKRKRLVLCLLEAADM KCFFPVLSCLAVLGVVSAQRQV TVQEGPLYRTEGSHITIWCNVS GYQGPSEQNFQWSIYLPSSPER EVQIVSTMDSSFPYAIYTQRVR GGKIFIERVQGNSTLLHITDLQA RDAGEYECHTPSTDKQYFGSYS AKMNLVVIPDSLQTTAMPQTL HRVEQDPLELTCEVASETIQHS HLSVAWLRQKVGEKPVEVISLS RDFMLHSSSEYAQRQSLGEVRL DKLGRTTFRLTIFH
4770	35138	A	4811	I	3728	MKCFFPVLSCLAVLGVVSAQR QVTVQEGPLYRTEGSHITIWCN VSGYQGPSEQNFQWSIYLPSSP EREVQIVSTMDSSFPYAIYTQR VRGGKIFIERVQGNSTLLHITDL QARDAGEYECHTPSTDKQYFG SYSAKMNLVVIPDSLQTTAMPQ TLHRVEQDPLELTCEVASETIQ HSHLSVAWLRQKVGEKPVEVIS LSRDFMLHSSSEYAQRQSLGEV RLDKLGRTTFRLTIFHLQPSDQ GEFYCEAAEWIQDP
4771	35139	A	4812	897	1217	ANRKASTMRWYVRPFCSGLST LSREKLYPLRMLRMRASTGPLS GALGMYSTASPRLMFSGCSISS AR*LFRSSAVTMPPAAWMSFTI WAVRLPL*KAWAPSMASVS

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SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
1	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
į				sequence		
4772	35140	Α	4813	1	1507	MWNSKINHDHDDQFLKTYYVL
'''	33110					ESSISVGVASNASVILEGDDLRF
				ŀ		SCSVRMAGRPQGRFSVIWQLV
			:			DRQNCRSNIMWLDRDGTVQPG
						SSYWERSSFGSVQMEQVQPNSF
	:					SLGIFNSRKEDEGQYECHVTEW
						VRVVDGEWQIVGERRASTPISII
						ALEMGFAVTAISRTPGVTYSDS
						FDLQCIIKPHYPAQVPVSVTWR
				j		FQPVGTVEFHDLVTFTRDGGV
						QWGDRSSSFRTRTATEKAESSN
						NVRLIISRASDTEAGKYQCVAE
						LWWKNYNNTWTQLAERTSNL
						LVIRVLQPDRMGVSARALRGEP
						SPSQMPQACCHGNTGALVIGIN
						EPESLPCLQTVTKLQVSKSKRT
						LTLVENKPIQLNCSVKSQTSQN
						SHFAVLWYVHKPSDANGKLIL
						KTTHNSAFEYGTYVEEEGLRAR
				-		LQFERHVSGDLFSLTVQRAEWL
						LSPNYAWYKLAEEVSGRTEVT
					:	VKQPGGSLGLGC\SVSGWAAEP
						VCTQGPRPCSWNTPLALTAAC
4773	35141	A	4814	1	627	
4774	35142	Α	4815	166	435	
4775	35143	В	4816	1	240	
4776	35144	Α	4817	1	288	VVPASSPGAASEPRRRRCLQP
						EKSVSPGSGGGHRDPPKARPPR
						PPSAPKP*RPRPFS*LARSLCFPA
						AGCAYGVGVGGAGGGRAGLR
						QPVPAVAE
4777	35145	Α	4818	206	1041	VSVGSAPSDMPAAATRW/CNSR
						CQHGQRLSAR/PVAGPGT/RTSS
						FPWLAPRNVVVPGSLEMPGTA
						GPQRGSHSSPQGSSALLSFS/CV
						HNVESKEPVCLLLPSMASRLLT
						PRGTCRPVPGCLQHPLSLPPVL
						VGTQSPSQAPKSAKRPSPQPQL
						GWLQLHPGGSGLCLLPAPAGS
						MECAALESVPSQLGIGAPGPCW
						AQAGVQGWSNVATSSPSGPSA
						QGQPRAPPCLVCGPAQCGTSGS
						RSRATGLRPSCLECQVWWSPQ
						CGANPGDTTSSQDAGTRCSKV
4778	35146	Α	4819	2	487	
L		1		L	<u> </u>	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ł	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4779	35147	Α	4820	152	758	RGARGPEWDGQLDGPGSQMC
						GMASAPRLRRSGVRAAAGHPV
						QQPRLLDGLSLTDESDSCVSFEP
		-				PSHPSSSSGPVTALSRGWNFLK
						AFPLMGETQERER/VLTHFSRR
						YCQCNPD/AQPSEDGIH/TLTCA
						LMLLNTDLHGHVIGEVTGETN
						RTDKQQNRTEQSKRQANADKH
İ						NEHANNEQQHEHDKDNDTRTR
ŀ						DTEMRHAKRDNA
4780	35148	Α	4821	100	220	
4781	35149	Α	4822	100	1588	CLKISTGYTGSMAEEASGNLTI
İ						MAEGEGEAGTSTHGQSRRKRE
						KEEVLHPFKQPDLVRSHYRENS
l						KEEICPHDSVTSCQTPPPTLGITI
						RHEIAASICHVADTAGVAGDA
						MMRKTDLIPPFIGLTLQGGRSA
						DMGNQKKWRKGSENLATLVA
						MASERFHWQYSGNSTIQPKQK
						QHGMITRLVTAPVCPGLVQPSL
						CAVGLRNIMSLSSHTCALQNTK
						QAWEGYDDCHSQRVSQPGEN
						MTKLSFLNLLNQMPSWGEGKM
						VPADFSGSCWEGTGTGSPAGSQ
						VSFLNRHNDGQLAGDSPGLPPP
						PCPRGLQGAEQRSALRRHKAD
						VHRDQWLVISFSKKLRTSARK
						WYEAGPFSGVPRVLPAAAQGG
		İ				TDFPFSPSEEPFQPSPFSMVQAH
		İ				TVRSKPQFHFPAAGENCLNRSE
						LSRNQGLQARVGPLAIMSSDPS
						GDFVLYTPTILGFVGLEVLVSR
						EGTFLWRYSTCP\LNYKLWLPP
						GYLGLLVPKDQKVRAVFLGM
4782	35150	Α	4823	1329	1881	AHLEAAGWSSSVQPGPSPRVLG
						RAEEHGQQSGRAKEIGHKIPKE
						TFDELTAELETISSK\CLYLAKK
						NQVIQQELLSMKEVQQKCKKL
						EEVNKILEQEVVNLKTHEKNM
						VEFGDVEECKLQLEERAGQEIE
						KLEEINLQRACLSAKKVRAEGE
			1			LEPRRHHNHTLGFTEGPWGLM
						SVITGAYDLIR
4783	35151	A	4824	176	308	LGNLTSPFFN*KHRSTKGFTTLS
						TVALLLTGHVGIKTSNCHPGY
						TVALLLTGHVGIKTSNCHPGY

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4784	35152	Α	4825	1467	1890	DWFDELTAELETISSK\CLYLAK KNQVIQQELLSMKEVQQKCKK LEEVNKILEQEVVNLKTHEKN MVEFGDVEECKLQLEERAGQEI EKLEEINLQRACLSAKKVRAEG ELEPRRHHNHTLGFTEGPWGL MSVITGAYDLIR
4785	35153	В	4826	1	753	
4786	35154	A	4827	225	410	
4787	35155	Α	4828	225	410	
4788	35156	Α	4829	225	411	
4789	35157	A	4831	141	453	GGVRGVQQKETCAFKVLESIG KLG\LALSVAGGAENSALYNVD AGHRAVIFDRFRGEQDIVVGEG THFLIPWVQK/PPIIFDCRSRPRN\ VPVITGSKDLQNVNITLRILF\RP VASQLPRIFTSIGEDYDERVLPSI TT\ENLKSVVAPFDAGELITQRE LVSRQV\SDDL\TERAATFGLIL DDVSLTHLTFGKEFTEAVEAKQ VAQQEAQRARFVVEKAEQQKK AAIISAEGDSKAAELIANSLATA GDGLIELRKLEAAEDIAYQLSR SRNITYLPTG\QSVLLQLPQ QPLPEPLELRPKAL*LTPSQIFSA
4790	35138	A	4831	194	455	*RLKTAARLPRKPWAVLPQGFT DTPPHFSQAQISSSSVTYLSIILV KTHVLCSPEFHRTGVV
4791	35159	A	4832	1	86	PCESYLEHL*WSCRLLRRLGSV SLQLLS
4792	35160	С	4833	191	263	
4793	35161	В	4834	1	741	
4794	35162	Α	4835	3	96	
4795	35163	A	4836	1	301	QEQQKMNTLQGPVSFKDVAVD FTQEEWRQLDPDEKITYGDVM LENYSHLVSLAYEVATSCTSEIL KPSNLPKSFFFSH*QDMISPSQT SSLSWSRERSCG
4796	35164	Α	4837	3	273	VLHFISAGNTFAHQQEHSPRKG PNNLSKRKLLPAV\GPRVFHGE DRHL/LFSTRKE*ARSLCYVQG GVQAPAAAFCSLLSLWGWAGC AFWC

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4797	35165	A	4838	3062	3688	RSVTRAPPRSTFPLAQAPSPRSH HIREPTPSPHLGPLRGSHRGPPS RPLPPALAALSLLETLQPRLLHP HLHL\LHDQGRACVRAYAHPS YLWRCHALGVWVES/ARLRPS ALPPAHSPPLAAPLLSSSVQPPA PALEPPPRPPAAGAAARCLAAQ HHHPSRLGVRQPPLAVGALGST WPPQAPAPPPELSALEQDRVGA QPPPPPSQGA
4798	35166	Α	4839	1	197	
4799	35167	С	4840	26	358	
4800	35168	A	4841	84	433	PASAPLGLSATVSACFQEQQKM NTLQGPVSFKDVAVDFTQEEW RQLDPDEKITYGDVMLENYSH LVSLAYEVATSCTSEILKPSNLP KSFFFSH*QDMISPSQTSSLSWS RERSCG
4801	35169	A	4842		372	VQWLFMQICLGGKHQQLRHQG VQEKMLPLEGS*VIEAGSPTLIG ETLSLEIINYIIGAGLESVHTHKS RQLGGWTRPGSHFQQVLIPAHT PMNLEPSQLSLEPRLVPAQQSL PSAPTPSKALSQ
4802	35170	В	4843	7	267	
4803	35171	В	4844	213	3895	
4804	35172	С	4845	141	359	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	lik .	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4805	35173	A	4846	831	3017	 RRRLHYSGSTRPWRTRAMLPR
4803	33173	A	4040	031	3017	ł.
]						TLSLRPHLSLSLHLPLPCLLLHL\
						LPTHLPPGPKHQPTVREL/ARRC
						RLPRACHPAQNLLEQQPLRPSH
	į.					LQRH/LSLLPPLGLSQSGPPG/LA
						PQPLL*LQTPDHFAGADPCSIHG
				:		TSH*HQGTSNPSGRDGYQTPSH
ŀ				:		ICP\SPAPKQSFLFGTQNTSPSSP
						AAPAASSAPPMFKPIFTAPPKSE
						KEGPTPPGPSVTATAPSSSSLPT
						TTSTTAPTFQPVFSSMGPPASVP
						LPAPFFKQTTTPATAPTTTAPLF
						TGLASATSAVAPITSASPSTDSA
l				:		SKPAFGFGINSVSSSSVSTTTST
	1					ATAASQPFLFGAPQASAASFTP
						AMGSIFQFGKPPALPTTTVTTF
						SQSLHTAVPTATSSSAADFSGF
						GSTLATSAPATSSQPTLTFSNTS
				.		TPTFNIPFGSSAKSPLPSYPGAN
						PQPAFGAAEGQPPGAAKPALAP
						SFGSSFTFGNSAAPAPATAPTPA
						PASTIKIVPAHVPTPIQPTFGGAT
						HSAFGLKATASAFGAPASSQPA
						FGGSTAVFSFGAATSSGFGATT
ļ				Ì		QTASSGSSSSVFGSTTPSPFTFG
						GSAAPAGSGSFGINVATPGSSA
						TTGAFSFGAGQSGSTATSTPFT
						GGLGQNALGTTGQSTPFAFNV
						GSTTESKPVFGGTATPTFGQNT
						PAPGVGTSGSSLSFGASSAPAQ
						GFVGVAPFGNTFAHQQEHSPR
		ļ				KGPNNLSKRKLLPAVRAQGPPR
4806	35174	Α	4847	9	935	IPCFGCAMPMYQTRRQENDLR
						TASIAV*RRKQQDDH*KQRRW
						QNIQRKGPKRYIVIAGNSQSHQ
ĺ						PMIFSMLRKLPKVTCRDVLPEIR
						AICIEEIGCWMQSYSTSFLTDSY
						LKYIGWTLHDKHREVRVKCVK
						ALKGLYGNRDLTARLELFTGRF
			:		1	KDWMVSMIMDREYSVAVEAV\
						RLLILILK\NMEGVLMDVDCES
						VYPI\V*ASN*ALASAVGEFLYW
						KLFYPECEIRTMGGREQRQSPG
						AQRTFFQLLLSFFVESKSHFVTQ
						GGGSGQFSAHRNLCLPGSGNFH
		<u> </u>				VSASRVAGIAGAPPHTWLIYVF
4807	35175	Α	4848	1	1749	
4808	35176	В	4849	282	1227	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4809	35177	A	4850	221	907	FEDSFNRGVKKKAAKRPLKTTP VAKYPKKGSQAVHRHSRKQSE PPANDIFNAAKAAKSDMQHRE VRVKCVKALKGLYGNRDLTAR LELFTGRFKDWM\VSMIMDRE YSVAVEAVRLLILILKNMEGVL MDVDCESVYPIVLFYPECEIRT MGGREQRQSPGAQRTFFQLLLS FFVESKLHDHAAYLVDNLWDC AGTQLKDWEGLTSLLLEKDQS TCHMEPGPGTFHLLG
4810	35178	В	4851	1	2361	
4811	35179	В	4852	1	878	
4812	35180	Α	4853		313	MPHTPRCLFRPQASCLVHFLEQ QNKLLETKLQFFQNRECCKSNL EPLFEGY\TLRREAECMEANSG RLASELNHVQEVLEGYKKKYE EEVALKATAENEFVALKK
4813	35181	Α	4854	188	354	
4814	35182	A	4855	405	647	LPLISDQVSYSPPLMSKSLASLY QESY*TILARIPLPLISPFVNSLST DLFILLIGNTSIAIFAEFGVSAISL PSCNSLE
4815	35183	A	4856	1	585	MSSGEGKRTWLGSLWGGVRVP VPPPICGLRVAVTSAGTSSFSLA PKPTRIRGERGKGQGRG/SSGN GGRGRAGASAGGSRGEVVSVA NDGRWRGRGRGVGCPRSSKRE DNRFAKHGRASGKAWEPHPPS QALRALFGPIRRRGRAAETRIV YWIKDRQLTNRDSTILELQKVL KTCCAQSMKIFCCLWNFVYKQ
4816	35184	C	4857	12	468	
4817	35185	A	4858		1156	MAFRKKIITAGCIDHVLSVSDS QQMQANLTQRTEAEIRKQTQIA PTFPAHSLPDKYLGFALLNRKII GTFLVFNMCKRNHEDKNEAFT VLREKKVFNLEFCTLQNYSSEV KEKERQRFSDKQTLREFATRKP ALKEMLKVLQRYKERKHIRKR NVKYSADHTTDHTANQADPNQ QEPHVPGHAPDNKTIRQKQNQ HKDNTTRKHNQMTSDTSRTNT KVSGQPRGKQTRRHERQQKA RERGQHRQK/RDAQTREHTQR QRQARAEE\QVIENKREEKRTS ERTNDSTRREGTRRPRRTEKGE RQEEGDRATATHRKDHDRQQP HGAPHCSRSSRHGAPTGPLYCC SPLPGTPSCTICTVQYGRLWPH MATEQLNMAGPNQDFQDVVW

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	1 .
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
4818	35186	Α	4859	1	1580	MAPALLLLLASGAAACPLPCV
		İ				CQNLSESLSTLCAHRGLLFVPP
						NVDRRTVELRLADNFIQALGPP
						DFRNMTGLVDLTLSRNAITRIG
						ARAFGDLESLRSLHLDGNQAG
						GAGHPRALRGPVNLQHLILSGN
1		ļ				QLGRIAPGAFDDFLESLEDLDLS
						YNNLRQVPWAGIGAMPALHTL
						NLDHNLIDALPPGA\FAQLGQLS
						RLDLTSNRLATLAPDPLFSRGR
		1				DAEASPAPLVLSFSGNPLHCNC
						ELLWLRRLARPDDLETCASPPG
						LAGRYFWAVPEGEFSCEPPLIA
						RHTQRLWVLEGQRATLRCRAL
						GDPAPTMHWVGPDDRLVGNSS
						RARAFPNGTLEIGVTGAGDAG
		l				GYTCIATNPAGEATARVELRVL
		l				ALPHGGNSSAEGGRPGPRTSPP
						PLALLPRYNSSEDETLIYRIVPA
						SSHHFLLKHLVPGADYDLCLLA
				}		LSPAAGPSDLTATRLLGCAHFS
				ŀ		TLPASPLCHALQAHVLGGTLTV
						AVGGVLVAALLVFTVALLVRG
						RGAGNGRLPLKLSHVQSQTNA
4819	35187	A	4860	2	403	
4820	35188	Α	4861	87	442	PVFPWHPYSKDQKEESQSVLVI
						WQKTAGGGWSTWWILSFSPET
						EKVVKK/AHVDTL*NEMWDVT
						K*TAFKIPSCT*HQQHP*SFFRK
					,	SESQVIVFAGQGVKIH*YILLHS
4821	25190		4862	2	651	NVIHYMNEI
4821	35189	Α	4802	2	031	EFIKQWDADKFEHIQTLEGHHQ EIWCLAVSPSGDYVVSSSHDKS
						LRLWERTREPLILEEEREMERE
						AEYEESVAKEDQPA\VPGEIQG
						D\SYFT*KKTIETVK\AAERIMEA
						FELYREETAKMKEHKAICKAA
						GKEVPLPSNPILMAYGSISPSAY
				ł		VLEIFKGIKSSELEESLLVLPFSY
						VPDILKLFNEFIQLGSDVELICR
						CLFFLLRIHFGQITS
			L	L	<u> </u>	CELEPTINI PORTO CELE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4822	35190	Α	4863]	703	MGLTKQYLRYVASAVFGVIGS
			:			QKGNIVFVTLRGEKGRYVAVP
		1				ACEHVFIWDLRKGEKGLKQEV
		İ				TCLCPSPDGLHLAVGYEDGSIRI
		İ				FSLLSGEGNVTFNGHKAAITTL
						KYDQLGGRLASGSK/RQPGGIV
						FTESILAYSSACQDKQNHYWGS
	İ	ļ	:			SQ*CADFVIQLRQYCCSFSCS*F
		1				H*NMEQVYTAVYSHNDL*ICTL LILCTW**TGSHRNKAPKTMFP
						RCTEGLDRGNRPATHS
4823	35191	В	4864	1	2616	RCTEGEDRGNRFATHS
4824	35192	A	4865	3	674	PECTGRTLRSASQQHDTFYILA
1021	33172	ľ.	1005			CELNSLADDAQR\YDVP\NSCL
				1		YPPDLPWRPVMQAKKKELARR
					:	DDIEDGDSMISSATSDTGSAKR
	1					KSKKNIRKQRMKILFNVVLEAR
	1					EPGSGRRLCDLFMVKPSKKDYP
						DYYIIILEPMDLKIIEHNIRNDKY
					:	AGEEGMIEDMKLMFRNARHYN
		İ				EEGSQVYNDAHILEKLLKEKRK
						ELGPLPDDDDMASPKLKLSKA
						ASHFII
4825	35193	A	4866	3	425	
4826	35194	Α	4867	1	1115	LCSVPTLCLASSHLVSNLSTFKQ
		İ				\TSSFIRHVVATLLGLYLALFCF
						GWYALHFLVQSGISYCIMIIIGV
		İ				ENMHNYCFVFALGYLTVCQVT
		•				RVYIFDYGQYSADFSGPMMIIT
		l				QKITSLACEIHDGMFRKDEELT
		l				SSQRDLAVRRMPSLLEYLSYNC
						NFMGILAGPLCSYKDYITFIEGR
			!			SYHITQSGENGKEETQYERTEP
						SPNTAVVQKLLVCGLALLFHLT
						ICTTLPVEYNIDEHFQATASWPT
			•			KIIYLYISLLAARPKYYFAWTLG TAPRKWSPCPGPQCIQLRAADP
						SGPEPLLMLSSQERQPKAKPPK
						EAHADIAGTPREGEKTRTGGPG
		İ				KAFLRDPGASSSVELSRTQLWV
						LTPPPIAKAPCECLQ
4827	35195	В	4868	218	656	21111111111111111111111111111111111111
4828	35196	A	4869	1	180	
4829	35197	A	4870	12	199	GLFPLEPGANGI*GC/SGRIRAQ
						RWLPGRPKFTGESFIR*PPRVAK
1						ESGQLIWFGCVPTQISS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
4830	35198	ĪA	4871	1 1	5058	MKLRGVSLAAGLFLLALSLWG
,030	30.70	` `		ľ		QPAEAAACYGCSPGSKCDCSGI
						KGEKGERGFPGLEGHPGLPGFP
						GPEGPPGPRGQKGDDGIPGPPG
	}	ł				PKGIRGPPGLPGFPGTPGLPGMP
						GHDGAPGPQGIPGCNGTKGER
		}				GFPGSPGFPGLQGPPGPPGIPGM
						KGEPGSIIMSSLPGPKGNPGYPG
		}				PPGIQGLPGPTGIPGPIGPPGPPG
		-				LMGPPGPPGLPGPKGNMGLNF
						QGPKGEKGEQGLQGPPGPPGQI
		1				SEQKRPIDVEF
4831	35199	Α	4872	3	1646	EDEGRAKGHHTWQQTRENESQ
		ĺ		Ì		AKGETPYKTIRFRETYYHKNSM
						GETTPMIQLSPTGSLPQHVGIM
						GATIQDKIWVTPGLPGFPGTPG
	:					LPVKRGFPGSPGFPGLQGPPVIP
						GPTGIPGPIGPPGPPGLMVTPGP
						PGLPGPKVNMGLNFQGPKGEK
1						VKQGLQGPPGPPGQISEQKRPID
				İ		VEFQKGDQVIPGDRGPPGPPGI
						RGPPVTPGGEKGEKGEQGEPG
Ì						KRVKPGKDGENGQPGIPVMPG
						DPGYPGEPGRDG/EKGNAVMG
ŀ						PPGPPGFPGERGQKGDEGPPGIS
						IPGPPGLDGQPGAPGLPGPPGPA
						GPHIPPSNKGDTCFNCIGTGISG
•						PPGQPGLPGLPGPPG\IPGAPGA
				ł		PGFPGSKGEPGDILTFPGMKGD
		ŀ				KGELGSPGAPGLPGLPGTPGQD
						GLPGLPGPKGEPVRITFKGERGP
]		PGNPGLPGLPGNIGPMGPPGFG
						PPGPVGEKGIQGVAGNPGQPGI
		}				PGNKGDPGQTITQPGKPGLPGN
						PGRDGDVGLPGIPGLPGQPGLP
		1				GIPGSKGEPGIPGIGLPGPPGPK
4832	35200	Α	4873	110	256	CSGTYRCYSF/HSRDPYLWSAPS
		1				DPLE/LVVTGPARQYYTKGNLV
1000	25201	 	1071		10000	RIASGL
4833	35201	Α	4874	2	2888	LSDPCSSRWDERSLSQRSRSWS
						YNGYYSDLSTARHSGHHKKRR
						KEKKVKHKKKGKKQKHCRRH
						KQTKKRRILIPSDIESSKSSTRR
						MKSSCDRERSSRSSSLSSHHSSK
						RDWSKSDKDVQSSLTHSSRDSY
	1					RSKSHSQSYSRGSSRSRTASKSS
						SHSRSRSKSRSSSKSGHRKRAS
						KSPRKTASQLSENKPVKTEPLR
					ľ	ATMAQNENVVVQPVVAENIPV
						IPLSDSPPPSRWKPGQKPWKPS
				L		YERIQEMKAKTTHL

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4834	35202	A	4875	350	986	LWGGGSARLLRLPRGPCLPPAS PSSPSSSSELS*SCSLGPSSSSSSS SSSCSLCPSSSSSSSSP*LPPSSSS SSSRPLPPPSSSSSSSSSSSSSSSSSSSSSSS
4835	35203	A	4876	1	496	SQSYS\RGSSRSRTASKSSSHSRS RSKSRSSSKSAVLASTARCAGS RKQCVPLPIRLCQLQSRSLQPS MVPWRWGLTGPRGIALPPAAR PGPLHEAPRWGPPPRRAPRPWP PGARPGRRRRRAAASPAPSSCG PRASGAAGGGRGAPIGARASA GAAVWTPISTT
4836	35204	Α	4877	3	4578	TLAVFVPTLAGFSVALGGPAW GRRRRSVSGVGVWLQWQCFLF CSRGPAQAGGQPALAATSVAM GAQDRPQCHFDIEINREPVGRI MFQLFSDICPKTCKNFLCLCSG EKGLGKTTGKKLCYKGSTFHR VVKNFMIQGGDFSEGNGKGGE SIYGGYFKDENFILKHDRAFLLS MANRGKHTNGSQFFITTKPAPH LDGVHVVFGLVISGFEVIEQIEN LKTDAASRPYADVRVIDCGVL ATKSIKDVFEKKRKKPT
4837	35205	Α	4878	1	1689	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
4838	35206	Α	4879	1	1604	MGISCPGSSKMLNSRALPVLKL
						TQGALRDCKERANYTNFNIYK
						AVESQGKFVCVLYNGQQSPEY
						VMIPCTSVSPTGQEVPEDRGMR
						MVMKNKPNKPSKMQSLKSSKH
						LTNSVIVIAVGIFFSVTLQARDP
		j				QLDDAIEQLRGVCIRAWEKITS
						GGEQYPSFSAIKQGPKEPYIDFI
		ļ				ARLQESLKKMIADSAAQDIVLQ
						LLAFDNANPDCQAALRPIRGKA
						HLVDCIKACDDIGDGSSNGKAS
		1		:		YFGSKSKVFQTSYTSAQKAELV
		İ				AVIEVLTAFDMPINVISDSSYM
						VYSTQLIENAQLRFHTDEQLMT
		ł			,	LFTQLQTAFRSTMHPFYITHIRA
		İ				R\HTPLPGPLTEGNQMADCLVA
!						NAISNARHFHNLTHVNASGLK
						HRYSITWKEAKNIIQRCPTCQM
						VHSSSFRGGVNPRGLEPNSLWQ
						MDVTHIPLFGRLAYVHKRRRIR
						GGGGGEGEGEGEEEEEE
ŀ		ł				EEEEEEEEEEEEEQEQEQEE
•						QEQEEQEQEERRRKDTE
		<u> </u>				ELSYSLPLNRTQPCWHPDFGLP
4839	35207	Α	4880	1	1146	
4840	35208	Α	4881	1	1577	MGQVWALVHSTLETFHTDEEE
		1		:		GEYNEVTEQVCLPAKAGSAAV
		1				DLCCTKAVSLLPGESPQKVPTG
						AGGPLPAGMTGLLLGRSSLNIK
·						AVQVQTGVTDSDYNGEIQIVTS
						TSVPWKAKPGDHIAQLLIVPKK
						F\EGLKEPLQVERQSSCQGLGY/
		1			:	PFLMAAIVKPPEPIPLKWLTDKP
		1				IWTEQWPLSKEKLEALEDLITQ
						QLKKGHIAPTFSPWNSPVFIIKK
		1				KSAEQDCEWFVFTILAVNNLQL
						KPAKRFHWKVLPQGPNQQPIWI
						PSRYLKPYHKPDAKEEIPEGSQ
						GFPVAAMSRLTLRRTPTVTSNT
		1				HRTQPPTWGQIEKLPQMAEENL
		ŀ		,	1	RKAGQPVTISNWILPRITKFKPI
		1	1			EGAENVFTDGSSNGKASYSGSK
			}			GPLTEGNQMADRLVAKVISNA
		1	1			RHFHNLTHVNASGLKRRYSIT
		1				WKEAKAIIQRCPTCQVMLSAAE
		1				QHLQKSAAKTEAEKLVWWRD
						PITKSREIGKIITWGRGYACVSP
}						GPNQQPIWIPSKHLKPYHKPDA
1	<u> </u>	1	1	1		GEKISGGNCGPHPHPRLQPCPD

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
4841	35209	ĪA.	4882	1	420	MTNISTTLKDLKDAGMIDPIKS
'' '	33207	`	1.502			TFNL/PIWIPQKPDGSWKTTANN
		İ				/R/KLNQATISIAAIVLDVASLPG
ŀ		1				KARSYHGFSSHAGRVIDSNNQK
						DIEGCTYILNVAGKNMFGIQLR
						RSVIFPCPNLMYTDNCSSHGPR
			}			GPSDYDSDP
4842	35210	A	4883	213	687	PEHILPGAQEGPSC/PLEPELE/V
4042	133210		7003	213	007	*NMSSWNTLSESCKMHGDLKY
	1		<u> </u>	1		LFVITAKDQQ/VSIAAST/*GFC
						WNQ*VSASPSQAPGMPLVLGR
						RLWQGSRLGGLSISLSDDCHGA
						HNHSFKRVTTTLQVFSWTVPQ
						AEPAMDCIHTA/DSPRQRLP*QA
						HESVAEVIIT
4843	35211	A	4884	1	648	MSSWNPLPESFKKHGYLLYLFV
4043	33211	^	4004		040	ITAKDQQ/VSITASTWGFYWNH\
						SQGKNPEQMEASQKPVAGLWA
						WWLNISLSAASHGAQNHPFKR
					Į.	VTTTLQSQGWNKDLWIMRELL
						CNNRPEQLLQETATIKAPIAPAD
					İ	PWSQEVIYNFCKAVSSVSCMPF
						NIHFNSNIPPESSGDWRMQQPC
						KVEQSILGAQRGGGGCRPGSLR
						ELFPCLLAEPNMEEEVAALRA
4844	35212	A	4885	3	597	GTLEPAAEWSVLLGV\HSQDGP
4844	33212	l ^A	4003	3	397	LDGAHTRAVAAIVVPANYSQV
				1		ELGADLALLRLASPASLGPAV
						WPVCLPRASHRFVHGTACWAT
				1		GWGDVQEADPLPLPWVLQEVE
		İ				LRLLGEATCQCLYSQPGPFNLT
						LQILPGMLCAGYPEGRRDTCQP
						SPEPGPMCIS/VSHIQRLRDAVS
			-			HTWQRGTEPARLGSWERRQRQ
						QQQGPV
4845	35213	В	4886	387	552	QQQ01 V
4846	35214	A	4887	20	612	
4847	35215	A	4888	792	959	TGLSSSQNPKATKSIPWHNRLL
		-		· · -		PNVDSQVQQNSQAIIYTN*GNS
						ESQYPFSKMDT
4848	35216	A	4889	268	500	<
4849	35217	A	4890	1	529	
4850	35218	В	4891	259	1672	
4851	35219	A	4892	480	657	,
4852	35220	A	4893	2	377	WCDFSSSLHRSVQALRTFCVTL
					1	SGAFVRDCTTVLSSTTCEPSTD
						V/CWAMNIVTSVLLLYGSEEEA
-						FWLLVALC\ERMLPDYYNTRV
						VVIRTYDQRYEQCVDLYDGLP
						TTLFTTDDWHVWPDLTGLP
L			L	<u> </u>	L	LIE TIDDWITY WIDEIGEF

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	1			sequence		
4853	35221	Α	4894	1	251	TEARKAAAVMPKTAVLAAERP
						KKAWGVLIPKKSTKRTPKAFA
						GGGCW/TKVGWSKVAKKVPKP
						EASKPKKAAPKTRRYKSTLKTK
4854	35222	Α	4895	283	427	PSVSKESPLNSPASQAGSATFAT
		ŀ				TPLTPPK/PAATSSVAAAALLPA
						LPR
4855	35223	Α	4896	111	1095	SRGWSRLPVPCCAPALLSPWAV
						NGIRRRGAGDGTRRGGSGCAG
						AHSAGLVASGPESWIPGPVPPG
						AEPPRRGDSGLARCPAPGHPSR
						ASPQRPAEGFDALAAPSRVPGD
						SRQLSRNNTRSRCEGRGGKTPP
						AWAGVHRHGPPPAAPPGRARV
	Ī					HGHRLRGGLGPGGGKRGLPGC
						CPRAADPRGRP/GETSRSGERGP
						PGRSAPLSLQPRHRTGLET/RQP
						SPLARSRPGLGPRPSARRPSRPA
						PPPPPPELHRGAPQRGRVASQT
						RPGARARPAADTHSSLKSPASQ
						AGSAAFATAPLTPQKPQLRAP
						WQPLHSFRPCRGVTSRLRTAPP
4856	35224	В	4897	243	452	
4857	35225	Α	4898	3	353	RRYLSPKYIKMFVLDEADEMLS
						RGFKDQIYDIFQKLNSNTQVVL
						LSATMPSDVLEVTKKFMRDPIR
	İ	İ				ILVKKEELTLEGIRQFYINVERE
	1			1		EWKLDTLCD\LSAMHGDMDQK
						ERDVIMR
4858	35226	Α	4899	3	410	NLQEWKLDTLCDLYETLTITQA
						VIFINTRRKVDWLTEKMHARDF
						TVSAMVCLPAASLLWVCPSEV
						SYLKPGFLEPRCLPGLLHICFLF
						QHGDMDQKERDVIMREFRSGS
						SRVLITTDLLVSRGN**QRQKG
						GSKVI
4859	35227	A	4900	1	235	MLSYVLEVTKKFMRDPIWILVK
						KEELILESIHQFCINVEQEKWKL
						DTLRDLYETLTITQA\VIFTNTR
						RKCCSKVREI
4860	35228	A	4901	3	48	NSNVEREEWKLDTLCDLYETL
7000	33440	``	1701	ľ		TITQAVIFINTRRKVDWLTEKM
						HARDFTVSAMHGDMDQKERD
						VIMREFRSGSSRVLITTDLLVSR
						GN**QRGTRGVEAGHTM
<u> </u>	<u> </u>	Ь	<u></u>	J	L	ON QROTROVERONIM

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4861	35229	Α	4902	2	879	AQFPTRLDLKATQALVLAPTRE
						LAQHIQKVVMALGDYMGASC
						HACIGGTNVRAEVQKLQMEAP
			i			HIIVGTPGRVFDMLNRRYLSPK
						YIKMFVLDEADEMLSRGFKDQI
				ł		YDIFQKLNSNTQVVLLSATMPS
						DVLEVTKKFMRDPIRILVKKEE
		ŀ	:			LTLEGIRQFYINVEREEWKLDT
		ŀ				LCDLYETLTITQAVIFINTRRKV
						DWLTEKMHARDFTVSAMHGD
		l				MDQKERDVIMREFRSGSSRVLI
						TTDLLARGIDVQQVSLVINYDL
		1				PTNR\ENYIHRIGRGGRFGRKGV
						AINMVTEEDK
4862	35230	Α	4903	1	1764	MASLEEGIYSLKINSKRSSYNS
						MNPQFTAGSEGQNETEEPNKFQ
						SRKGRPQDSGSSLSLSVLQEILG
						DDIPKVKEEAVISSQSDLGECPY
						CGERPARNATIFAPQKEKESAP
						EMSSSCDKRVTVNPPEKFSEGR
					}	PKTQNTLICEKCSQPSNFLDDY
						NPHVIIQKRLSNQSRVICEKSSP
						PLNVPDNYNSHSVIQKHKEKM
						AIERPSSGSDWSDVGGTTVIFSE
						EKPFSLCLPVVSEPPYYATDYT
						TFPPHYSPWHDYTSSWFSSTKS
						SCYPSLGSSSNTLQAGKSSCSSS
						SRSNNNIFQGERSSHQSFLSDYS
						TSFPVSSENTSRDLKMTEEGRS
						KNSSLFYYSRNVEAEAKEERVY
						QEETLGHPYGGRASSFPPRTIW
						QPEQPGFIDTHCHLDLLYSRLPF
						KGTFTKFRKIYSNTFPKEFQGCI
						SCFCNPQNLSDNLWEDQLKDD
		ŀ				LVWGVFGCRPHFAHYYNNYQE
						RSILKALRHPKAVAFGEMGLD
						YSSKCTTHIPEQQKVFKRLLRL
						AVSLK\KPMMIHCREADEDLLG
						ILKKYVPSDHKMYQHCFTGSYP
						VIKPLLSCFPNLYVGFMAILTYS
				<u> </u>		SAEQARETVKK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4863	35231	ĪA	4904	1125	820	VLAPNYHITFRNLSSVSSPVTSA
1005		[]				EPVLQAGQARELSFSSLHPASPL
						LVPSTPALTLHAFNALVLFPSFP
						LEKAAKRERPKSKIQITARVGK
		1				MRRTWSPHTTISAGGSVNGAST
		1				LKNSLVWEAKPADTTNPTPPTT
•		1				HPNPGPTPPTNTNRHPPPHHPLT
		Ì				NTPAPPHQPPATTP/SPQTQTPN
						YCPNTDHTTPTHDTPNPVPPTR
						RTPPNRTGKRTNTSQNYRLVSP
						APKIHWEA
4864	35232	A	4905	407	757	7 THIT WELL
4865	35233	A	4906	5	426	GKTSTKSQNASSPPKDHNSSPA
						REQN/WIENEFNELTEVGFRRIT
						SLEKNINDLMELKNTARELREA
						YTSINSQINQVEERISEIEDQLNE
	İ					IKHEDKIREKRMKRNQQSLQEI
		1				WDYVKRPNLRLIGVPESDIENG
						TKLENTL
4866	35234	l _A	4907	1	2267	MTGQFQDVVREEARPLPNRIM
1000	323 .	`				RLRFNHFATECSWDHLYVYDG
	1					DSIYAPLVAAFRCEVENRYQGN
						PLRGTCYYTLLIDYQFTFSLSQE
						DDRYYTAINFVATPDEETPPIRG
	1					RQTAHTGELQLTSGGYPSGMK
	1			•		LPEEGTGERNRITINKKDVHTET
						PSKGHQHQRPKVDKSTKIKKN
						QRKKAENSKNKNASSSPEDHN
						SSPAREQNWMENEFDELTEVG
		}				FRRWVITNSSKLKKHVLTQCKE
	İ	1				AKNLEKRLGQLLTRITSLEKNV
		1				KELMELKNKAOELCEAYTSINS
		1				QIDQAEESISEIEDQLNEIKHED
	1	l				KIREKR\IKRNKQSLQEIWDYVK
						RPNLRLIGVPESDGENGTKLEN
	1					TLQDIVQDNFPSLARQANIQIQE
						ILKTPQRYFSRRATPRNIIITFTK
						VEIKGKMLRAVREKDHSAIKLE
						LRIKKLTONCTTMWKLNNLLL
	İ					NEYWVHNEIKAEINKFFETNEK
		l				KDTTYQDLWDTAKAVFKGKFI
	1					ALNAHRRKWERSRIDTLTSQLK
	-	1				ELEKQEQTNSKASKRQEITKIR
						AELKEIETQKTLQKINESRSWFF
						EKINKIDRLLARPIKKKREKNQI
	1					DTIKNDKGDITTEPTEMQTTIRE
						YYKHLYANKLENLEEMDKFLD
	1					
		1				TYTLPRLNQEEVESLNRPTTSSE IEAIINSLPTKKSPGPDGFTAKL
						YQRYKEEVVPLLLKLFQTVGK
						EGPLPNSFYWANIILIPKPGRHT
L	<u> </u>		<u> </u>	<u> </u>		EUTETINST TWAINILLIPKPUKHT

SEQ ID	SEQ ID NO:	1	SEQ ID NO:	į.		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
1	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
4867	35235	С	4908	605	760	
4868	35236	Α	4909	94	400	
4869	35237	A	4910	1	1538	MPIQMQYPQYKLVENKTQPPVI
						YQYWPPADLQYRPPPEVKYRP
						QVVCPVPNSTAPYQQPTAVVF
		ŀ				NPTAPPSGQGIIAQNTDLVEWF
						FLPHSTIKTFTLYLDQMATLIGQ
						ARLRIIKLCESDSDKIIVPLNKEQ
						VRQAFINSGAWQIGLADFVGII
		l				DNHYPKTKIFQFLKLTTWILPKI
						TRHKPLENALTVFTDGSSNGKA
		1				AYTRPKERLIETQYHWAQRAE
				}		LVAVITVL\QDFNQSINIVSDSA
						YVVQATKDVETALIKYSIDDQL
					}	NQLFNMLQQTVRKRNFPFYVT
						HIRAHTNLPEPLTKANEQVDLL
						VSSAFLEAQELHALTHVNATGL
		ŀ				KNKFDITWKQAKNIVQHYTQC
				}		QIPHLPTQEAGVNPREQHFTGK
				İ		KNSPHEGKLIWWKDKKNKTW
						EIGKVITWERGFACVSQGENQL
						PGWIPTRNLKFYNEPLGDAKKS
	į				ł	ASAETKNPQLSIIDSPGKAPGCL
						MPTTQNWLVEVPTVSATSKFT
						YHMVYPPAPKRQRPARTGHND
						DGSFVKKGDM
4870	35238	Α	4911	1	759	FRMVIRCLPQRELDCDRGNWT
						ATSAELTGIKWRRYNFGGHGD
						CGPIISVPAQDDPILLSFIRCLQA
						NLLCVWRRDV*PDC*ELWIFW
						WGDEPNLVGV\YIMNCRLWKK
						DSGKMAFPMNVGRC/FFKAIHN
						LLERCLMDKNFVRIGKWFVRP
		ļ				YEKDEKPVNKSEHLSCAF\TFFL
		ļ.				\HGESNVCTSVEIAQHQPIYLIN
		ŀ				EEHIHMAQSSPAPFQVLVSPYG
	ė					LNGTLTGQAYKMSDPATRKLIE
		ŀ	_			EWQYFYPMVLKKKKK
4871	35239	A	4912	1	539	MRHPCDGNSLYFDHISVNMLA
				1		VMLYYGFGRVSIGKTGLRGVQ
						RHLWPLPLDASSTKTVSSSHSD
						KGEIFSPFPCSLINCNNFEMRNT
		1				TKCLMDKNFVRIGKWFVRPYE
						KDEKPVNKRVTSQEIWLFDEY\
						GVSLPCFAVERCALLPLCLLPR
				1		LMISISSTSRQPQYISRRGLKAKI
						swvv
4872	35240	Α	4913	198	620	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4873	35241	IA	4914	63	1267	RNRILRMEFFDWFWAFWSWLL
						NMIRSQNAKDSSSNSMENTDSP
						WCELFRELCKINALDVPDSLLV
						RGNEFSASVHNTFDHMWRTKE
						RYEAGWLLLSSADRVMKENDE
	ŀ					LRDSVSWLQKQILSLKSAKIAL
	[SGSLISYRERAEIVEKQTQTLIM
						RVADLQQKGYINSPALCHNLIW
						RDLDHFSFPYDITLFHYIDDIML
	ŀ					IGSSEQEVANTLDILKGQRFVLT
						VIDTYSRYWFASPECNASA/KT
						TIHGFTECIIHYHGIPHSIASYQG
						THFMAEEPGW\SKNQGVEVTPF
						TITPSDPVATFLLPVPMTLRSAG
						LEVLVPEGGTLPPGDTTMIPLN
						WMLRLPPGHFGLLLPLSQQAK
						KGVTVLAVVIDPDYQDEISLLF
						HNGAIISDGNGVILIGLLQWFST
4874	35242	Α	4915	2	2210	
4875	35243	Α	4916	1	2036	MDTFLDTYTLPRLNQEEVESLN
			}			RPITGAEIVAIINSLPTKKSPGPD
		ŀ				GFTAEFYQRYKEELVPFLLQLF
						QSIEKEGILPNSFYEASIILIPKPG
						RETTKKENFRPISLMNIDAKIVN
						KILAKRIQQHIKKLIHHDQVGFI
						PGMQGWLNTCKSINVIQHINRA
						KDKNHMIISIDAEKAFDKIQQPF
						MLKTLKKLGTDGTYFKIVRAIY
						DKPTANIILNGQKLEAFPLKTGT
						RQGCPLSPFLFNIVLEVLARAIR
	-					QKEIKGIQLGKEEVILSLFADD
		1				MIVYLENPIVSAQNLLKLISNFS
					ŀ	KVSGYKTNVQKSQAFLYTNNI
						QTGSQIMSDLPFTIASKRIKYLG
						IQLTRDMKDLFKNYKPLLTEM/
						KTKKWKNIPCS/WELEKTTLKFI
						/WNRKRARIAKSILSQKNKAGG ITLPDFKLYYKATVTKTAWYW
İ						
						DQNRDIDQWNRTEPSEITPHIY NYLIFDKPEKNKQWGKESLFN
	}					KWCWENWLAICRKLKLDPFLT
	-					PYTKINSRWIKDLNLRPKTIKTL
	1					EENLGITIQDIGMGKDFMSKTP
	1					KAMATQAKIDKWELIKLKSFCT
						AKETIIRVNRQPTKWEKIFTTYS
	1	Ì				SDKGLISRIYNELKQIYKKKKQP
						HQKDFAYHFTVFVFYFGAFLLE
	1					AAATSLHDLHCNTTITG/HATPE
						**PV*HKRSSLTITYLCRSG*IIW
	1					WCDPHTHS
1876	35244	1_	4917	 	1350	WODITING
4876	35244	Α	4917	1	1359	WCDPHIHS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4877	35245	В	4918	1	948	
4878	35246	Α	4919	1	981	
4879	35247	Α	4920	1	3747	
4880	35248	Α	4921	1	1875	
4881	35249	A	4922	1	2238	
4882	35250	Α	4923	1	3924	
4883	35251	Α	4924	1	1068	
4884	35252	В	4925	62	389	
4885	35253	В	4926	1	3663	
4886	35254	Α	4927	1	1474	MAVSTTVRVDMPCCAVMLYC
						WGRSGKEQQPDEALKGKGFTV
•			:			CFEGLRFGQKPGSYFSLDQIGH
		l				VEIQMITPPWRGTTLSTKRGGR
						KRGKTESDREHNLAPPKGRDTP
		1				KKEGAHVRGGPLSGTPPTEKK
						GKARSPRYGDTHRGEGLNKTH
						RAPONFSESPEPTINKIDRPLAR
						LIKKKREKNQIDTIKNDKGDITT
						NPTEIQTTIREYYKHLYANKLE
•	}			!		NLEETOKFLOTYTLPRLNQEEV
ŀ			İ			ESLNRPITGSEIEAIINSLP\TKKS\
•		1	:			PGTDGFTAKFYQRYKEELVPFL
		1				LKLFQSIEKEGILPNSFYEASIILI
		1				PKPGRDTQKKENFRPISLMNID
		l				AKIVSKILANRIQVHIKKLIHHD
						QVGFIPGMQGCVGSSARAIGQE
						KEIKGIQLGKEEAKLSLFADDTI
			:			ECLENPIVSAQNLFKLISNFSKV
						SGYKIHVPKSVSFLYTNNIQAES
						QIKNAIPFTIASKRIKYLGIQLTL
						EVKELYNKNYKTLLKEIRDE
4887	35255	Α	4928	419	1002	CYQRWKVLKKSSGPLARL/IKK
						REKNQIDAIKNDKGDITTDPTEI
						QTTIREYYKHLYINKLENLEET
						DKFLDTYTLPRLNQEEVESLNR
						PITGSEIEAIINSLQTKKSPGPDG
						FTAKLYQRYKEELVIYRFNAIPI
						KLPMTFFTELEKTTLKFIWNQK
						RIRIAKSILSQKNKAGGITLPDF
						KLYYKATVTKTAWYW

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				Sequence		
4888	35256	Α	4929	1	2502	MFFETNENKDTTYQNLWDTFK
						AVCRGKFIALNTHKRKQERSKI
						DTLTSQLKEPEKQEQTHSKASR
						RQEITKIRAELKEIETQKTLQKIL
						QKINESRSWFFEEINKIDRLLAR
		l				LIKKKREKNQIDAIKNDKGDITT
		l				DRTEIQTTIREYYKHLYANKLE
		•				NLEEMDKFLDTYNLPRLNQEE
						VESL\HRLITGSEIEAIINSLPTKK
						SPGPDGFTAKFYQRYKEELVPF
					:	LLKLFQSIEKEGILPNSFYEANII
						LIPKPGTDTTKKENFRPISLMNI
						DVKILNKILANRIQQQIKKLIHH
]	DQVGFIPGMQGWFNIHKSINVI
<u>.</u>		1				QHINRTKDKNHMIISIDAEKAF
<u> </u>						DKIQQPFMLKSLNKLVLEVLAR
					1	AIRQEKEMKGIQLGKEEVKLSL
						FADDMIVYLENPIISAQDLLKLI
	-					SNFSKVSGYEINVQKSQASLYT
	}					NNRQTESQIMSELPFTIASKRIK
						YLGIQLTRDVKDLFKENYKPLL
				1		NEIKEDTNKWKNIPCSWVGRIN
	}					IVKMAILPKVIYRFNAILIKLPM
				ľ		TFFTELEK\TTLKFIRNQKRACIA
						KSILSKKE\KAGGIMLPEFKL*Y
						KATVTKTVWYWCQNRYIDQW
						NRTEPSEIIPHIYNHLIFDKPDKN
						KKWGKDSLFNKWCWEHWLAI
	-					CRKLKLDPFLTPYTKINSRWIK
						DLNVRPKTIKTLAGHLGNTIRDI
						GLGKDFMTKTPKAMATKAKID
1000	10000	<u> </u>	1020		1107	KWDLIKLKSFCTAKETAIRVNR
4889	35257	A	4930	1	4187	MGDFNTPLSTLDRSMRQKVNK
						DSQELNSALHQADLTDICRTLH
						PKSTEYTFFAAPHHTYSKIDHIV
	}					GSKALFSKCKRTEIITNCLSDHS
						AIKLELRIKKLTQNRSTTWKLN
				-		NLLLNDYWVHNEMKAEIKMFF ETNENKDTTYQNLWDTFKAVY
1						1
1						RGKFIAFKAVCRGKFIALNAQN
						RKQERSKIDTLTSQLKELEKQE
						QTHSKASRRQEITKIRAELKEIE
						TQKTLQK\ISESRSWFFEKTNKI DRPLARLIKKKR
	l	<u> </u>	<u>i </u>	L	<u> </u>	DILLAKLIKKK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
4890	35258	Α	4931	1	1818	MSSRGPCTLETLINPSPSVADDI
						PHLRPKPVYITTTRDNENIYSTK
ļ						IPYMAARVVFIKWIVTFFLEKK
ļ						YLTATQNTKNGVDVLPKIIQTV
İ						GGGAVQERAPELDGGGPTEQD
						KSHSNSSTLSDRRLSNSSLCSIE
						EEHRMVYEMVQRILLSTRGYV
						NFVNEVFHQAFLLPSCEIAVTR
						KVVQVYRKWILQDKPVFMEEP
						DRKDVAQEDAEKLGFSETDSK
						EASSESSGHKRSSSWGTH\NSFT
						SAMSRGCVTEEG/TIQMLKAGV
				-		QALLQPVLCVFPLIPLPPLCNQA
						ARPAVVSIPLGDLLPWLSVLGG
						PADQWHLSYLKPEGSSAYACIL
						EALESAGKGLWKVSAVGRKP
						WTRGRSDRKGTVGHQVQFLFF
						KDEGRVDSRILTLMVAWIRPN\
						LCVYISRELWDDFLGVLS\SLTE
						WEELINEWANIMDSLTAVLAR
						TVYGVEMTNLPLDKLSEQKEK
						KQRGKGCVLDPQKGTTVGRSF
						SLSWRSHPDVTEPMRFRSATTS
						GAPGVEKARNIVRQKATAKRS
						QSISNCVHLSEALPATKSVPLLL
						HTLYKNKALKAALGQPVGKAD
						QLVVAGSPSVCCVEDQWGRSN
						LEAMVLWLYCIPPDGDVLYL
4891	35259	С	4932	96	278	
4892	35260	Α	4933]]	261	KLRLPPGHFGLLLLLSQQAKKG
						VTVLAGVIDPDYHNEISLLLHN
						GGKKKTRPAEMLAEGKGNTE
			100		0.05	WVGEEVSHQYQL*PHDQLQKR
4893	35261	С	4934	36	385	
4894	35262	A	4935	207	2349	OLA EDDLIWIDGED AD A COLUCT
4895	35263	Α	4936	207	561	QLAEPPHWIPGTRKDAQCLVGL
			1			FGFWRQHIPHLGGLLWPIYRVT
						QKAASFEWGPE*EKALQQIQPS
						VQAALPLGP*DLADPMVLEVS
				1		MADRCAGWSLWQAPIGELQQ
	<u> </u>	<u> </u>				KPLGFWSKALP

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4896	35264	A	4937		1375	MSTTQLDTMGAESLCNMETVQ VEDEGIQKLKEIRMVEWITPFRP THPSCEGPEDIPLTNALQNTFVR AAPASLKSPVVALLLSPVGYRT HAVVISPVPECRFGIDILSSWQN PHIGSLAGRAHQKQFVGSWQG QQYAFTVLPQGHINSLALCHNL IWRDLDHFLLSQGITLVHYTDD IMLIGSNLADPMVLEVSMADRC AGWSLWQAPIGELQQKPLGFW SKALPSSADNYSPFERQFFACY WALVETECLPTGHQLTMQPEL PIMNWVLSDPSSHKWVHGQSG HGGRDSGYSWTQQHGLPLTKG DLSMTTAECQIFQQQRPILSPRY GTIPWGDQPATWRQKWQRFVL TGIDTYSRYEFAYPACHASTKT TIHGLMEFLIHHHGIPHSIASDQ GTHLMAKEVRQWAHAHGIHW SYHVPHHPEVAGLIERWNE\GL
4897	35265	A	4938	1	324	LKSQLQHQLVNRLRRELQCWL INCLRNCKTYQA/RKPLWFYNT SLKFFLNKP/MLADVVFEIQGTT VPAHRAILV/ARCEVMAAMFN AGIFQAMCLLICAE/MYQVSRL QHICELFII/TQLQSMPSRELASM NL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	1	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
4898	35266	I IA	4939	1108	2304	L VAAAYPPPASPPWSPARSLAGA
1070	33200	l'`	1,,3,		2501	APAPLCVGPAAVEAREGDAAG
		ŀ				DERIAGELAARGPREAVSRCFS
		1				PSRRPALGFEIMSIHIVALGNEG
						DTFHQDNRPSGLIRTYLGRSPL
		ŀ				VSGDESSLLLNAASTVARPVFT
						EYQASAFGNVKLVVHDCPVW
		1				DIFDSDWYTSRNLIGGADIIVIK
						YNVNDKFSFHEVKDNYIPVIKR
						ALNSVPVIIAAVGTRQNEELPCT
						l `
						CPLCTSDRGSCVSTTEGIQLAKE
		1				LGATYLELHSLDDFYIGKYFGG
						VLEYFMIQALNQKTSEKMKKR
						KMSNSFHGIRPPQLEQPEKMPV
						LKAEASHYNSDLNNLLFCCQC
						VDVVFYNPDLKKVVEAHKIVL
						CAVSHVFMLLFNVKSPTDIQDS
		ŀ				SIIRTTQDLFAINRDTAFPGASH ESSGNPPLRVIVKDALFCSCLSD
		l				ILRFIYSGAFQWEELEEDIRKKL
:						KDSGDVSNVIEKVKCILKTPGKI
						NCLRNCKTYQARKPLWFYNTS
						LKFFLNKPMLADVVF\EIQGYG
		l				QCPAHRAIL\VAPCEVM\AAMF
						NGNLHGKAKSVLIPVYGVSKET
		l				FLSFLEYLYTDSCCPA\GIFQAM
						CLLICAEMYQVSRLQHICELFII
						TQLQSMPSRGTGHPWNLDIS/V
						DLL*KGPSFHHSWIAFSTWAYF
						HFHCYLTYLIFQFKRPGIFRDLF
						S/EGRNGSFLFGKRHRWGPSNM
1000	25267	_	4940	1	3117	LLGRQLAGITGKYYFTSPGNVG
4899	35267 35268	B A	4940	1	1162	MRVCARACVRTRTRMCVYAH
4900	33208	^	4741	1	1102	TCVCVRTYAYVRVRVRVHVRA
						RVRVCAYARTRVRNSLSILPFIQ
		ŀ				
						LTLATPIHHIHQEEFNIRGIVPVL RRVKPDLAIGIDITPSCDTPDLH
						DYSEVRINQGVGITCLNYHGRG
						-
						TLAGLITPPRLIRMLEQTALEHN
						IPVQREVAPGVITETGYIQLFLP
		l				GWEIGFSPLALLLAFLCSTSPGF
						GDPDGLGVIAYQDTVRPNAAT
-		1				AISELNALAVKGVILTGDNPRA
						AAAIAGELGLEFKAGLLPEDKV
						KAVTELNQHAPLAMVGDGIND
						APAMKAAAIGIAMGSGTDVAL
		1				ETADAALTHNHLRGLVQMIEL
						ARATHANIRQNITIALGLKGIFL
						VTTL\MTGLWLAVLADTGATV
	1					LVTANALRLLRRR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4901	35269	A	4942	1889	3387	QWAIKQFKHAEKENPYLFGGG
,,,,,,	133203	,	1.5.2	1.003		HLRAIFR*SGFQ*RQLHTMHWG
						AS/QQAEHP/LAQPSTRAQLRTPI
						P/TADS/QRALVGSGMKPRLNG
						ERVLICAAGKHPADAFTGLINE
	1					LESAGQTVVLVVRNDDVLGVI
						ALQDTLRADAATAISELNALGV
	}				ļ	KGVILTGDNPRAAAAIAGELGL
						EFKAGLLPEDKVKAVTELNQH
	İ					APLAMVGDGINDAPAMKAAAI
						GIAMGSGTDVALETADAALTH
					<u> </u>	NHLRGLVQMIELARATHANIRQ
						NITIALGLKGIFLVTTLLGMTGL
						WLAVLADTGATVLFFPRLFEAR
		1			l e	NSQVGNGETNQTRFRLCAAPG
ļ						GTFIADFTAGTGRRARPRRNRR
		1				RVVVSFDFHQNMRRFLMEIVA
						ARFVVSKVAAHFRTFHYGGVIF
						ISRENVIWRGFESIFDHLEQRFR
			,			LLFTIDNPVGVKNFVAAVLGVR
						LGKHIQFDVVRVTTKLCESILQI
						VNFIFRQSQAKTQVSVDQRLTA
						LPQQINAGNRSRLMVGKQLLCI
4902	35270	В	4943	1	2104	
4903	35271	С	4944	1	1215	
4904	35272	Α	4945	3	268	YEFNRPYPEIQRSGIS\KLLEPLL
i		1				FAATSDSQLSKTEISSIKINSETV
						PVYQLRYNGNNALMFATYQD
		1				KMLVFSSTDMLFKDDQQDTEA
4905	35273	В	4946	1	1725	
4906	35274	Α	4947	1	1437	
4907	35275	Α	4948	2	736	QCPYPYIAATGRILVVTHLICLA
		1				PQIVENQVTYPLTTTMLSVPGA
						KTVRGFSQFGDSYVYVIFEDGT
						DPYWARSRVLEYLNQVQGMLP
						AGVSAELGPDATGVGWIYEYA
		1				LVDRSGKHDLADLRSLQDWIL
						KYELETIPCVSEVASVGGVVKE
						YQVVIDPQRLAQYGISLAEVKS
1						ALDASNQEAGGS\SI\ELAEAEY
						MGGTCSAHGRALSRGVLWCRF
					-	SAIVRGPPRGMFCGAGAAPNLI
4908	35276	A	4949	531	632	
4909	35277	Α	4950	532	609	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
4910	35278	A	4951	11	525	L LGVEATA\IGDFGK\AAEKLPVP\
710	33276	()]	323	VELAQACAHAVMKSGIDLAVS
						YCMQVNHGFAQPLEFLLGGLD
						KVPVLLVFINGVATPLPGFQRT
						RMLGEAIGRFTSTPQYAGDTVN
						DEDISNTIRALFATGNFEDVRVL
ŀ			ļ			RDGELRKFPTCESDDNFRPVTF
						SVNYVIRHDWVGIRRVVPENE
4911	35279	A	4952	1	2769	STATE VIOLET VIEWS
4912	35280	A	4953	1	681	
4913	35281	Α	4954	1	1230	
4914	35282	Α	4955	1	774	MLRTSDRPPYQYPKRCIPLSVL
				•		EVLRCARRWSLRPYKQLARAYI
						/ERYRHDR\DIEREKRGRLNQEI
						RGLVEQTNASLLNENANKDSK
·						AHERGDIHYHDLDYSPFFPMFN
						CMLIDLKGMLTQGFKMGNAEI
						EPPKSISTATAVTAQIIGSVHIYG
						GTTINRIDEVLAPFVTASYNKH
						RKTAEEWNIPDAAGYANSRTIK
						ECYDAFQSLEYEVNTLHTANV
						EGLTGDDGIAVEALKKNSQKPR
						MSASHNCELVSLRGRQYT
4915	35283	Α	4956	1	1278	
4916	35284	Α	4957	461	1002	SLSTKGYAAAKWCSTACAGS/T
		1				LLLIVLLWVFIPKGFFPGQDNGI
						IQGTLQAPQSISFANMAQLQRQ
		1				VADVILQDPAVQSLTSFVGVDG
	į	1				TNPSLNSARLQINLKPLDERDD
		1				RVQKVIARLHTAVDKVPGVDL
	ļ	1				FLQPTQDLTIDTQGSRTQYQFT
		1				LHATSLDALNTWVPQLMEKI\Q
						QLPKLS
4917	35285	Α	4958	1	291	MNPVDRPLLDIGLTRLEFLRISG
	:	1				KGLA\GLTIAPALLSLLGCKQED
	ļ					IDSGTVGLINTPKGVLVTQRAR
						CTGCHRCEISCTNFNDGSVGTF
		<u> </u>				FSRIKPH
4918	35286	Α	4959	1	306	MKNFEVLQPLQNSLSGLPLWV
						SERILQQINQLTHYEPVIGIMGK
						TGAGKSSLCNALFAGEVSPVSD
						VAGCTRDPLRFRLQIGEHFMTI
						VDL\PGVGESGVRD
4919	35287	Α	4960	2666	2775	
4920	35288	Α	4961	9	308	GDLDGSRHRAFLCHLPVELKTA
				1]	LMFPVTLTRALMETAY/ATAVS
				1		A/NFRTESRGAH/SRFDFPD/RDD
						ENWL/CHSLYLPESESMTRRSV
						NMNPKLRPAFPPKIRT
4921	35289	Α	4962	1	1132	1

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	i .	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4922	35290	A	4963	1170	1352	RNLPDAAGQSCAGEIRLYRQQT QRSRLQRWLHLACFRRCNLHR
						RELDATA*QHVRSCRWR
4923	35291	A	4964	1	312	MRKLTALFVASTLALGAANLA
						HAADTTTAAPADAKPMMHHK
						GKFGPHQDMMFKDLNLTDAQ
						KQQIREIMKG\QRDQMKRPPLE
						ERRAMHDIIASDTFDKVKAEA
4924	35292	Α	4965	3	205	KDHIKGVLEAQNSLSTQVSLFF
						*SFFG*NSNMRNLIQIYRSL/CTK
						AI*LMNFKRHKNQLVINYLSRS
4925	35293	Α	4966	1	915	MTGVATARARGLRRPLSVGQV
						PPFARLWKGVQTVERHMLANI
						GTGSQHPGCGFQLINQNSVIVF
		1				AAGEVNGFTSGDVQCLKMRSG
						DMNDIQRRQRLLPDGNKFGEIQ
				1		TTIREYYKHLYANKLEDLEEM
		i				DKFLDTYTLPRLNQEEVESLNR
						PITGSEIEAVVNSLPTKKSPGPE
						RLTAEFYQRYKEELVPFLLKLF
		1				QTTEKEGLLPNSFYEASVLLIPK
						PGKDTTKKENFRPISLMNIDVKI
		1				LNKILANRIQQHIKKLIHHNQV
						GFIIGVQGWFNICKSTNVIYHV
						NRTNDKNH\IISIDTEKAFD
4926	35294	Α	4967	1	1338	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	İ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
4927	35295	Α	4968	<u> </u>	1928	PPLVIHRQTGSGEDLQQTPTDL
17,21	33273	ļ'``	1700	l '	1,720	QLRVLTIRRKTNKQKGHPHQNP
				ļ		ISSRRHEITKIRAELKETETQKTL
						QKKSMNPRSWFF\ERINKIDRLL
		i				k -
		1				ARLIKKKREKNQID\AIKNDK\W
						DITHRIPTEIETII\REY\YKHLYT
		l				NKL\EN\LEEMDKFLDTYTLP\R
		ŀ		1		LN\QEEVESL\HRLITGSEIEAITN
]						SLT\TKKSPGPDGF\T\AEFYQRY
						KEELVPFLLKLFQSIEKEGILPNS
						F\YEASIILISKPGRDTTKKENFR
	1					PISLMNIDAKILNKILANQIQQHI
						KKLIHHHQVGFIPGMQGWFNIL
		l				KSINVIHHINRTKDKNHMIISIEA
						EKAFDKIQQPFMLKTLNKLGID
						GTYLTYLKIIRAVYEKPTANIIL
						NGQKLEAFPLKTGTRQGCPLSP
						LLFNVVL\EVLVGAIRQEKEIKG
						IQLGKEDVKLSLFADDMIVYLE
						NPIVSAQNLLKLISNFSEVSGYK
l						INVQKSQAFLYTNNRQTESQIM
						SELPFTIASKRIKYLGIQLTRDV
						KDLFKENYKPLLNEIKEDTNK
	ĺ					WKNIPCSWIGRINIVKMAILPKT
İ						LNQKFSYWFRVNKHYIHQRTFP
	ŀ					
		1				LKETEFNTIATLYNGASP/RTAP
1						KSTGTNGHQASGLPRF*RIAFCS
	l					ALVKSKRKLYQGYLPGQTDRR
		ļ				EEGVSWCPGGP
4928	35296	A	4969	2	237	QPVEDTWLSTPAAPMVDSLIAR
						VRVMARGNAITLPVCGRDVK/F
				}		TLEVLRGDSVEKTSRVW\SGIET
		<u> </u>				YQELVTEDALDDL
4929	35297	Α	4970	1	1752	
4930	35298	Α	4971	1	723	
4931	35299	Α	4972	1	1701	
4932	35300	Α	4973	1	1446	
4933	35301	Α	4974	1	403	MKLMETLNQCINAGHEMTKAI
						AIAQFNDDSPEARKITRRWRIG
						EAADLVGVSSQAIRDAEKAGR
						LPHPDMEIRGRVEQRVGYTIEQI
						NHMRDVFGTRLRRAEDVFPPVI
						GVAAHKG\LALKGLRVLLVEV
						RHRV
4934	35302	Α	4975	1	1374	
				<u> </u>	 	L

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
4935	35303	I A	4976	11	2259	MKLMETLNQCINAGHEMTKAI
4933	33303	^	7770	'	2237	AIAQFNDDSPEARKITRRWRIG
						EAADLVGVSSQAIRDAEKAGR
					:	LPHPDMEIRGRVEQRVGYTIEQI
			:			NHMRDVFGTRLRRAEDVFPPVI
						GVAAHKGGVYKTSVSVHLAQ DLAL/KGLRVLLVEVHKRWS*N
						DEGYRHCTV***\HPEARKITRR
						WRIGEAADLVGVSSQAIRDAEK
						AGRLPHPDMEIRGRVEQR\LVIQ
						LNKLIICVMCLVRDCDVLKTYF
						HR*SGLLPIKVAFTKP/HV\$VHL
						AQDLALKGLRVLLVEEPPSAPN
						LGIGTINVVCAADVLIVPTPAEL
						FDYTSALQFFDMLRDLLKNVD
Ì	ŀ					LKGFEPDVRILLTKYSNSNGSQ
						SPWMEEQIRDAWGSMVLKNV
						VRETDEVGKAAPMVDSLIARV
						GVMARGNAITLPVCGRDVKFT
:						LEVLRGDSVEKTSRCDNLKTCH
						TSHGSVMAETAVINHKKRKNS
						PRIVQSNDLTEAAYSLSRDQKR
						MLYLFVDQIRKSDGTLQEHDGI
						CEIHVAKYAEIFGLTSAEASKDI
						RQALKSFAGKEVVFYRPEEDA
						GDEKGYESFPWFIKRAHSPSRG
				İ		LYSVHINPYLIPFFIGLQNRFTQF
						RLSETKEITNPYAMRLYESLCQ
						YHPLTIEKVMKPLITSNTVTDEI
						ERANVFKMNGKWYLFTDSRGS
						KMTIDGINSNDIYMLGYVSNSL
						TGPYKPLNKTGLVLQMGLDPN
4936	35304	В	4977	1	1744	
4937	35305	Α	4978	1	2367	
4938	35306	Α	4979	1	1215	
4939	35307	Α	4980	1	946	MKLMETLNQCINAGHEMTKAI
						AIAQFNDDSPEARKITRRWRIG
						EAADLVGVSSQAIRDAEKAGR
						LPHPDMEIRGRVEQRVGYTIEQI
	İ					NHMRDVFGTRLRRAEDVFPPVI
						GVAAHKGGVYKTSVSVHLAQ
			}			DLALKGLRVLLVEGNDPQGTA
						SMYHGWVPDLHIHAEDTLLPF
		1				YLGEKDDVTYAIKPTCWPGLDI
		1		1		IPSCLALHRIETELMGKFDEGKL
1						PTDPHLMLRLAIETVAHDYDVI
				[VIDSAPNLGIGTINVVCAADVLI
		1				VPTPAELFDYTSALQFFDMLRD
						LLKNVDLKGQAPCMK\LTMRS
1		1		1		SSSSAPSPWML
4940	35308	A	4981	1	1572	SOSSITI ST WINE
7240	00000	17.	7701	11	13/4	

SEQ ID NO:	SEQ ID NO: of peptide	1	SEQ ID NO:	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO.	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4941	35309	Α	4982	1	1617	
4942	35310	Α	4983	1	550	
4943	35311	Α	4984	1	3818	
4944	35312	Α	4985	607	3777	
4945	35313	Α	4986	671	3091	ARCQRVPGRPVPDLLRCLPRAT
						TPAPSPAPGLVVRRLAAPVPAA
						APVPAAAPVPAAAPVPAAAPVP
		1				AAAPVPAAAPVPVAAPVPVAA
		ļ				PVPAAAPVPAAAPVPAAAPVPA
		l				AAPVPAAAPVPVAAPVPAAAP
						VPRFQSPNQTASTSDSPHIGQLP
						HQTVPISDSFHTGQSPHRTDSTP
						DSPHTGQLPHRTVPTPD/ELPHR
						TVPTPDSFHTRQSPHQTASTPDS
						PHIGQLPHQTVPISDSFHTRQSP
		ļ				HQTASTQDSPHTGQIPHRTVPT
	j	ŀ				PDSFHTRQSPYRTASTPDSPHIG
						QLAHRTVPTPDSFHTGQSPHRT
						DSTPDSPHTRQLPHQTVPISDSF
		ļ				HTGQSPYRTASTPDSPHTRQLP
		•				HQTASTPDSPHTRQSPHRTVPT
		ŀ				PDSSPHQTVSTPDSPHTRQLPH
						QTVPTPDSFHTRQSPHQTVPTP
		ļ				DSPHTRQLPHQTASTPDSP\TPD
						SPHTGQSPHQFPHQTVPTADSP
		ŀ				HTRQLPHQTVPTADSPTPDSFH
						TRHSPHQTASTPDSPHIGQLPHR
						TVPTPDSFHTRQSPHQTASTPDT
						PHTRQLPQQTVPTADSPTPDSF
						HTRQSPHQTVPTPDSPHTRQSP
						HRTVPTPDSFHTRQYPQRTVPT\
						QTASTPGSPHTKQSPHQTVPTP
						DCPHTGQSPYQTVSTPDSPHTR
						HSPHQTASTPDSPHIGQLPHQT
						ASTLDSPHIGRHPRQMAPTPESP
						HARQLPCQMAPTPDSPTLDGPH
4946	35314	В	4987	1	891	
4947	35315	Α	4988	157	1990	
4948	35316	В	4989	1	2157	
4949	35317	Α	4990	1	378	MSIKGTYLKVIKAIYDKPTASSI
		}			1	LNGENLKAFLLRTGTRQGCPLY
		ŀ				KINVYKSVALLYTTNDQGENQI
						NNSTPYTTAPRKIKYIGIYLTKE
						VKDLYKENYKTLLKEVIDDT/R
				}		WKHIPCSWMGRINIV
4950	35318	Α	4991	1	392	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4951	35319	A	4992		1570	ETDFAFVGQAGVQWCDLSSPQ PLPPWVQAILLPQPPRQGCWSG IPKWSGISEDVQVWKNCKLDL AAATGKNCLGQGEEALLTQMR TGNQREVSDWWEEFVYLRSRN PLMVNSNYYMMDFLYVTPTPL QAARAGNAVHALLLYRHRLNR QEIPPVRGPQWVRDGGVVLWP LGHVGPGRQLTAHGFLQTLLM GMRPLCSAQYEKIFNTTRIPGV QKDYIRHLHDSQHVAVFHRGR FFRMGTHSRNSLLSPRALEQQF QRILDDPSPACPHEEHLAALTA APRGTWAQVRTSLKTQAAEAL EAVEGAAFFVSLDAEPA\GLTL EDPAASLDAYAHALLAGRGHD RWF\DK\SFTLIVFSNGKLGLSV EHSWADCPISGHMWEFTLATE CFQLGYSTDGHCKGHPDPTLPQ PQRLQWALPDHIHSSISLALRG AKILSENVDCHVVPFSLFGKSFI RRCHLSLDSFIQIALQLAHFRDR GQFCLTYESAMTRLFLEGRTET VRSCTREACNFVRAMEDKKKT
4952	35320	A	4993		1027	MPAKSPGQDVLGTDNIKIDVFK QTNQKSHTGEKPYECPECGKAF SEKSRLRKHQRTHTGEKPYKC DGCDKAFSAKSGLRIHQRTHTG EKPFECHECGKSFNYKSILIVHQ RTHTGEKPFECNECGKSFSHMS GLRNHRRTHTGERPYKCDECG KAFKLKSGLRKHHRTHTGEKP YKCNQCGKAFGQKS\QLRGHH RIHSGEKP\YKCNHCGEAFSQKS NLRVHHRTHTGEKPYQCEECG KTFRQKSNLRG\HQRTHSGEKP YECNECGKAFSEKSVLRKHQR THT\GEKP\YNCNQCGEAFSQKS NLRVHQRTHTGEKPYKCDKCG RTFSQKSSLREHQKAHPGD
4953	35321	A	4994	154	331	IPAAATCMGSLLGG*ETPGLWA RRSVKSRGLFPGLPSPSRASVRS LLLLPAWAAFLEGIVDTRPTAW RAFPWTLFLSVFCQFLDFPETSL DSQKLSLDTPSF

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	coden for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4954	35322	A	4995	657	1455	PEPKPAAVAPPPESPLPAGDSEG
"						CLQPGHLL\PASPRLPRPRQAAS
						LAHRPEKSAPGADCPSGRR\GR
						PWTCGGLSGPAAAPRPCPLAPA
ŀ						FAPAFLALGAQR\GPGPEQGGH
						ERQAERPGRRSRRTRAYSARIY
						LPVAAEWPMGPGPEPGAGGRR
						RGEVPGPGAQRSPARRRRGGPA
						AVSSGGRGVGGLRGAVLFQHP
						EQQQRPVGLAGLGAQQPAHV
						WRYGPLST/PPALGPPAAARGP\
						GRRLFLPFSFSPSAGAPGAASRP
						SPRRSP
4955	35323	Α	4996	ī	415	-
4956	35324	В	4997	8	211	
4957	35325	Α	4998	1	563	MDPNHEEIPDLPEKEFRRVKSK
1	•					NHMTISIDAEKAFDNIQHRFMI
						KTLSKIGIQGTHFN\IIKAIYDKP
						TASIILNGENLKAFPLRTGTSIVL
						EVLARAIRREKEIKGIQIGKEEV
						KLLLFADDMIVYLENPKDSYRK
						LLEQIKEFSKVSRYKINVQKSIA
	:					LLYANSDQAENQINNPTSFTIA
						AKNKIK
4958	35326	Α	4999	667	960	
4959	35327	Α	5000	317	1157	EFRMDPAIALVLALSLTVPKLT
						VPDSPLRLYRSFCTSACTSNLV
				:		VHFA*KEKWPDV*LYTDSWAV
	ì					ANGLAGWSGTWKKHDWKIDD
						NEIWGRGMWIDLSEWSKTVKIF
						VSHESAHHIT*KSSAEEDFNNQ
						VDRMIHSVDTTRPLSPATPVITQ
						WAHEQSGHGDRDGGYAWAQQ
						HGLPLTKADLAMVTAECPICQQ
						QRPTLSP*YGTIPRGDQPATWW
						QVDYIGPLSSWKGQRFVLTGID/
						TSGYEFAYPARSSANSTIRGLM
						ECLIHHHGIPHSIASNQLYS
4960	35328	Α	5001	2	15246	

SEQ ID	SEQ ID NO	: Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	<u> </u>	<u> Ш</u>				
4961	35329	A	5002		951	MALALNRYL/CLAVLPLITKCA PLFAG/TEHRAIMVDSMLHTVY R/LSRGRSLTKAQRDVIED/CLM SLC\RLLRRLVFDVPILNEFA/K MPLKLLTNHYERCWKY/YCLPT GWANFGVTSEEE/LHLTRKLFW GIFDSLAH/KKY/DPELYRMAM PCLCAIAG/ALPPDYVDASYSSK AE/KKATVDAEGNFDPRPVET/L NV/IIPEKLDSFINKFAEYT/HEK WAFDK/DKEIYRWPIKESLKAM I/AWEWTIEKAREDLDPREGYN PQPPDLSAV/TLSRELQAMAEQ LAENYHN/TWGRKKKQELEAK GGGTHPLLVP/YDTLTAKEKAR DREKAQSY
4962	35330	A	5003	273	478	Didition 1
4963	35331	A	5004	2	622	GAPRSLSEKERQLMGMINQLSS FREQLLHAHYEQKKLAASQIEK QRQHMKLGKQ\QQEQIARQQQ RLIQQQHKINLLQQQIQQLYAA QLAAMQVSPGGKLPGIPQGNL GAAVSPTSIHTDKSTNSPPPKSK TCCLLLWEDEHSASIPEVTNQQI HLHLASSITDHGGKVIWERCLE ESGKGVTGFPDWSPRWGRIPPE GSKEGFTKE
4964	35332	Α	5005	3	512	
4965	35333	A	5006		642	MHAQTYAHTDTRAQPCTSMEK QVSLKPTEEAHKKERKPQKPG KYICQYCSRPC/AKPSVLQKHIR SHTGYG\CEMYPHGLEMERIPG EEFEEPTEGESTDSEEETSATSG HPAELSPRPKQPLLSSGLYSSGS HSSSHERCSLSQSSTAQSLEDPP PFVEPSSEHPLSHKPEDTHTIKQ KLALRLSERKKHQVIDRLGSGI LAGLSWQLLPQGEAL
4966	35334	A	5007	341	1020	TLNRRQGCCLGSWLTRSNGILL FEISRPTYLPILLILFLVKR/HKL KMSVLNMVVRHDPGSTESAKA KEELIFRCRSRHFRASPLFSQHA AADKHKFQRFLTADMALAVTV YAPTTFPPSSLLLFKQKSNGMH NLIATGHLLLVDPNRMVIKRVV LSDPPFKICTKMAVVCYMFFNR EDVQWFKPVELRTKWSRRGHI QEPSGTHGRMKCSFDRKLKSQ NTELMNLYK

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4967	35335	A	5008	1	1847	EKAQALEDLAGFKDPAPGHTE
1,707	33333	['`	5000	'	047	ESMTHDKTTKIPCKSPPELVDT
			}			ATSSKKWPRACGQKVEVKEEL
						LAVGKLTQTSGETTHTNKEPEG
						EGKGMKAFKQPAKQKLDPENV
						TGSRRLPRVPKEKAQPLEDLAS
				,		FQELSQTPGHTEELANGADSFT
						SAPKQAPDSGKPLKISRRVLQA
						PKVEPVGDLVGTR/D/PVKSQSK
						SNTSLPPLPFKRGCGKDGSVTG
						TKRPHCMPAPEEIIEELPASKKQ
1						R/VVAPRVRGKSPEPLVIMKRSL
						RTSAKRIEPAEDLNSNNMKTNN
4968	35336	A	5009	9	1094	IRHEEESMTDDKTTIIPCKSSPEL
1,700						EDTATSSKRRPRTRAQKVEVKE
ļ						ELLAVGKLTQTSGESTHTDKEP
						VGEGK\GTKAFKQPAKRKLDA
	1					EDVIGSRRQPRAPKEKAQPLED
						LASFQELSQTPGHTEELANGAA
						DSFTSAPKQTPDSGKPLKISRRV
						LRAPKVEPVGDVVSTRDPVKS
						QSKSNTSLPPLPFKRGGGKDGS
						VTGT\KRLRCMPAPEEIVEELPT\
						SKKQRV\APRARGKSSEPVVIM
						K\RSL\RTSAKRIEPAEELNSND
						MKTNKEEHKLQDSVPENKGISL
						RSRRQNKTEAEQQITEVFVLAE
l						RIEINRNEKKPMKTSPEMDIQNP
	}					DDGARKPIPRDKVTENKRCLRS
						ARQNESSQP
4969	35337	Α	5010	1	459	MAKPWRALCILLRSLDSFLQLS
						ASLTCFGFMHNTYKNRPWGVL
		ŀ				RSSTERSPRKGTAGDMRAGGRF
						NAPLIGQDLLFKEHFASKALDT
						CTNIARSAKTQGGPVAAILLLL
						AFGPCIFNLRVKFVSSRIKAIKL
L						QM/RLTIYRGPLDRPAGPSTGL
4970	35338	В	5011	1	687	
4971	35339	С	5012	194	418	
4972	35340	Α	5013	17	396	
4973	35341	Α	5014	1	70	

NO: of peptide hod in USSN			Amino acid sequence (X=Unknown,
	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
sequence 09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	sequence		
4974 35342 A 5015	1	994	MGFVKVVKNKAYFKRYQVKF
			RRRREGKTDYYARKRLVIQDK
			NKYNTPKYRMIVRVTNRD\IICQ
			IAYARIEGIMIVCAAYAPELPK\
			YGVK\VGLTNYAAA\YCTGLLA
			GPAGFFHRFGMDKFYEG\QVEV
			TGDEYNVESIDGQPGCLHPAY
			WDARPLPRTSPLAIKVFWLP*R
			GA\VDGGLVLFPPQYPNRFPWL
			WIPESQGNLNARRYHREGTIHG
			PRIVAGLHAPTLMEEDEDA\YK
			KQFVRQYVKNS\VTPD\MMEE
			MYKKAHAAIR\ENPVYEKKPK
			KEV\KRKRWNR\PKMSLAQKK
			DRVAQKKASPL\RAQERAAEKL
			NOTIFYKDFSDIAN
4975 35343 C 5016	27	308	
4976 35344 C 5017	68	307	
4977 35345 A 5018	388	1485	
4978 35346 A 5019	3	1940	RDRMILLKMEQEIIDFIADNNN
			HYKKFPQMSSYQRMLVHRVA
			AYFGLDHNVDQTGKSVIINKTS
			STRIPEQRFCEHLKDEKGEESQ
			KRFILKRDNSSIDKEDNQSVCS
			QESLFVENSRLLEDSNICNETYK
			KRQLFRGNRDGSGRTSGSRQSS
			SENELKWSDHQRAWSSTDSDS
			SNRNLKPAMTKTASFGGITVLT
			RGDSTSSTRSTGKLSKAGSESSS
			SAGSSGSLSRTHPPLQSTPLVAG
			\VAAGSPGCVP\YPENGIGGQVA
			PSSTSYILLPLEAATGIPPGSILL
			NPHTGQPFVNPDGTPAIYNPPTS
			QQPLRSAMVGQSQQQPPQQQP
			SPQPQQQVQPPQPQMAGPLVT
			QSIPAKKELNAFRVRNQEQRDT
			GTLHEGSGKISRMVLGWEPLSP
			YAPDRDDVATQFGQMTLSRQS
			SGETPEPPSGPVYPSSLMPQPAQ
			QPSYVIASTGQQLPTGGFSGSGP
			PISQQVLQPPPSPQGFVQQPPPA
			QMPVYYYPSGQYPTSTTQQYR
			PMAPVQYNAQRSQQMPQAAQ
			QAVLEVLSPLVSTVMPPKLAVL
			VMAGFRLRLELSESVLLQALW
			WSDHLSSFSELLCRLPDVLPEPS
			LLPLEYDHQKDAQGQDYLTCM
			VLVTLLLESPTGIQEDDSNEISVI

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
4979	35347	IA	5020	3	1353	GDAYALVLRVTLYCGKCH\NE
777	33317	` `	3020			VVLAPIFEILSTESIQDQLPYSV
						MLMSMLAITEGRRSFSVSVENA
			Ì			CSNYVTTVQVKEVNQMHISPN
						SCNAIHPRDHILEISGTPTHTLO
						VEELEDAISQMSQTLLLLIEHDP
					-	VSRSPAQLRLDAWLSPHTQNA
						GHPHALSTLDTKDNLEGTLRRC
		1				SLRLSNSISKSPSPSSPKEPLLLS
						RDISCWESLCCSSTYSKQNFRSC
						DLIHGEVLGKGFFGQAIKVTHK
j						ATGKVLVMKELMRCDEENQK
			İ			TFLTEVKVTLPSNLLPPGHDLL
						QTGAREQTGILKTGGLSVPGGA
						GHPAACRAGGVGPHCEHAVRP
						DPRLTPLALAQPPAGGCSTASIA
						PLFSIPAVSRVFRASRGLAACLE
						AEQAGRTILTCEVCKLVVGLSL
						DSQQQLSSVLCGWLCLSCPQL
						WSLNRHPQHQHHLVEMQNLKS
						HPRPGETESAL
4980	35348	Α	5021	1	588	MEYPDGGSTLDLLEPGPLDETQ
						IITILREILKGLDYLHSEKKIHRG
						VKAANVLLSEHCEVKLVDFGM
						AGQLADTQTKRNTFVGTPFWI
						APENNPPTLEENYSKPLKEFVE
						ACLNKELSFRPTAKELLKHKFIL
		1				RDTKKTSYLTELIDRYKRWRA
						KQSQEDSSSEDSNSETDGQDS\G
4981	25240		5022	 	1200	GSDSGDWIFTIQEKDPKNL GPRASPPRADRRGPAVPSQEVA
4981	35349	Α	5022	1	1380	VLSAMAHSPVQSGLPGMQNLK
						ADPEELFTKLEKIGKGSFGEVF
j						KGIDNRTQKVVAIKIIDLEEAED
						EIEDIQQEITVLSQCDSPYVTKY
						YGSYLKDTKLWIIMEYLGGGS
						ALDLLEPGPLDETQIATILREILK
		ŀ				GLDYLHSEKKIHRDIKAANVLL
						SEHGEVKLADFGVAGQLTDTQI
		l				KRNTFVGTPFWMV\PEVIKQAG
		İ				YDSKADIWSLGITAIELARGEPP
						HSELHPMKVLFLIPKNNPPTLE
						GNYSKPLKEFVEACLNKEPSFIP
						TAKELLKHKFILRNAK\KTSYLT
						ELIDRYK\RWKAEQSHDDSSSE\
						DSDAE\TDGQASGGSDSGDWIF
						TIREKDPKNLENGALQPSDLDR
						NKMKDIPKRPFSQWLSTIISPLF
		1				AELKEKSQ\ACGGELGGPLKEL
						RGAIYLAEEACPGISDTMVAQL
			1			VQRLQRYSLSGGGTSSH
			1	1	1	

SEQ ID	1 -		SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
4982	35350	Α	5023	1	346	MAGEKVEKPDTKEKKPEAKKA
						DAGGKVQEGTGRYSRSAMYSR
						KAMYKRKYSAAKSKIEKKKEK
						VLATVTKPVGGDKNGGTQVF\
						QIITYSSYTQKVQLPKSTLKQRQ
						GPCPQGAL
4983	35351	Α	5024	67	599	SAAKSEMAGEKVEKPDTKEKK
						PEAKKVDAGGKVKKGNLKAK
		ł				KPKKGKPHCSRNPVLVRGIGRY
		}				SRSAMYSRKAMYK\RKYSAAK
						SKVEKKKKEKVLRNLLQNQVG
						GDKN\GGTR/VGLNFRKMPRI\Y
						PTE\DVP\RKLLSHGQKNPFSQH
		ŀ	,			V/REKLRASITPGTILI\ILTGRHR
						GKRVVFLN
4984	35352	Α	5025	101	202	PVPGTSSCTDRYHYKKH**KIN
						TKLVSKYILNM
4985	35353	Α	5026	2	562	LRTVPDLPGRRFRAMRTGQRR*
						PELPPDMNSLEQAEDLKAFERR
						LTEYIHCLQPATGRWRMLLIVV
						SVCTATGAWNWLIDPETQKVS
				-		FFTSLWNHPFFTISCITLIGLFFA
	1					GIHKRVVAPSIIAARCRTVLAEY
		}				NMSCDDTGKLILKPRPHVQ*QS
						SLIVMGLRIAFLRISDTAKSHKG
l						FLLRLDM

588

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	1	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
4986	35354	Α	5027	1	2090	MGAVFFMLFSWWITSGHIQLLC
						TRRRATVSTHQEFVVTVSPFSL
						GLVVSGCLDADTAMPGSAEEQ
						FQWQSQDGQKDIEDELTTGLEL
						VDSCIRSLQESGILDPQDYSTGE
						RVIHFGKTLGLSSEDYSIMEKG
						QMVKEVTVTPQVSVLKANETM
1						ASALCGQEGHDDPLLLTVQMS
						TGNHYIKDNSLFPWVKWPSLLS
		ļ				QSALQLNSKPEGSFQYPASYHS
						NQTLALGETTPSQLPARGTQAR
						ATGQSFSQGTTSRAGHLAGPEP
						APPPPPPPREPFAPSLGSAFHLPD
				1		APPAAAAAALYYSSSTLPAPPR
		ŀ				GGSPLAAPQGGSPTKLQRGGSA
						PEGATYAAPRGSSPKQSPSRLA
						KSYSTSSPINIVVSSAGLSPIRVT
						SPPTVQSTISSSPIHQLSSTIGTY
						ATLSPTKRLVHASEQYSKHSQE
						LYATATLQRPGSLAAGSRASYS
	ł					SQHGHLGPELRALQSPEHHIDPI
						YEDRVYQKPPMRSLSQSQGDP
						LPPAHTGTYRTSTAPSSPGVDS
						VPLQRTGSQHGPQNAAAATFQ
						RASYAAGPASNYADPYRQLQY
						CPSVESPYSKSGPALPPEGTLAR
]						SPSIDSIQKDPSKYPILCDGGNPF
						CALVYRPSSRERASSL/LKSMCY
						Q*NRLYPKFRRVLGSGFEGMQ
						RRFCASKGIKGRFCGSSLEQDC
						EHGGQHPCYPKLVPTLWNPHQ
						REGNQKNSERV
4987	35355	A	5028	123	3825	REGINQRINSERV
4988	35356	A	5029	1	593	MQVCSSGGRILRVSGTERQLRR
1700	33330	1.	302)	'	373	EGGRFCLGRGVRNLREGGALR
						APFTAPSLTALPPDIVDIKPAN\
						MEDLTEVTTASESHPHHCNLFV
						YSSSKGFLRLCDMRAAALCDK
						HSKHPCDDIGPIQIIQGHLRTSR
						GWMIILPALQSTFRCRMDEQNL
						RAFLAPRLCDFSTSQPQPGSPEQ
1000	25257	_	5020		004	PVMSGLISPWTKPSRSSVFSM
4989	35357	Α	5030	1	894	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	1	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
4990	35358	A	5031	<u> </u>	3115	I RQICSMASELGARDDGGCTELA
4990	33336	IA.	3031	1	3113	l `
						KPLYLQYLERALRLDHFLRQTS
	1					AIFNRNISSDDSEDGLDDSNPLL
						PQSGDPLIQVKEEPPNSLLGETS
						GAGSSGMLNTYSLNGVLQSES
		1				KCDKGNLYNFSKLKKSRKWLK
						SILLSDESSEADSQSEDDDEEEL
						NLSREELHNMLRLHKYKKLHQ
						NKYSKDKELQQYQYYSAGLLS
		1	1			TYDPFYEQQRHLLGPKKKKFK
						EEKKLKAKLKKVKKKRRRDEE
		<u> </u>				LSSEESPRRHHHQTKV
4991	35359	Α	5032	1	1522	MVAQASGQTIREAEVGGSLTA
		ŀ				QEVEATSILLSDESSEADSQSED
						DDEEELNLSREELHNMLRLHK
						YKKLHQNKYSKDKEWMSQLA
						ALNYYLAGGDEQKLLDLSYSPP
						QNRQPAMAKAHNNNNNNNNN
						NNNNHNKNPKRLLDERELLTY
						STATQSFDTRNTRSFDEDAKES
						RAAALRAANKSGTGFGESYSL
						ANPSIRAGEDIPQPTIFNGKLKG
						YQLKGMNWLANLYEQRLERL
	1					RFLQLEQEHTCPQEGRAERLRG
						QPLKLWALLHFIMPTLFDSHEE
						FNEWFSKDIESHAENKSAIDET
						KLKKVKKKRRRDEELSSEESPR
						RHHHQTKVFAKFSHDAPPPGT
						KKKHLSIEQLNARRRKVWLSIV
						KKELPKKTLYTQDAPFHVVITS
						NQLVVQDVKYFQRVKWQYMV
						LDEAOALKSSSSLTRSNARYPL
						VSSQRKIHDQQVTTDKSSKMDP
						KEEEYIPAKHEMENTKLQFYLS
						AIGEKQ\VRWKILLQFQCRNRL
						LLTGTPIQNTMAE
4992	35360	A	5033	1	974	MGFHHVARAGLELLTSGMCCY
1,,,,,	33300	`		1		FAQDLRPEQSIKASLQRIILRKY
		İ				EKCGHHNLQLKKGYKSVDEYK
					}	VHKGSYNGFNQCLTTTQSKIFQ
						CDKYVKDFHKFSNSNRHMTEK
						NPFKCKECGKSFCVLSHLTQHK
						RIHTTVNSYKLEECGKAFNVSS
1						TLSQHMRIHTGQKHYQCGEM\
						GIAFNKSSHLNTHKIIHTGEKSY
						KREECGKAFNISSHLTTHKIIHT
		1				GENAYKCKECGKAFNQSSTLT
						RHKIIHAGEKPYICEHCGRAFN
		1				QSSNLTKHKRIHTGDKPYKCEE
		1	1			CGKAFNVSSTLTQHKRIQQQQQ
		<u>L</u>				QTIKQTKNLLNNNNKKLWV

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
4993	35361	В	5034	22	684	
4994	35362	Α	5035	1	1338	
4995	35363	A	5036		596	MGMTDMGKSKKKSQGPPSFPP GGAPPPAIGAPAGRERPPGTGD PFNQRPCDNLKTCHTSHGSVM AETAVINHKKRKNSPRIVQSND LTEAAYSLSRDQKRMLYLFVD QIRKSDGTLQEHDGICEIHVAK YAEIFGLTSAEASKDIRQALKSF AGKEVVFYRPGEDAGHEKGYE SFPW/FIKRSPE*RRGKFPFTQM GQD
4996	35364	Α	5037	787	1692	
4997	35365	Α	5038	289	1338	
4998	35366	Α	5039	982	1347	
4999	35367	Α	5040	1	780	
5000	35368	В	5041	148	334	W 7
5001	35369	Α	5042	2	545	
5002	35370	Α	5043	1	732	
5003	35371	Α	5044	1	1110	
5004	35372	A	5045	1	1170	
5005	35373	Α	5046	1	870	
5006	35374	Α	5047	1	1254	
5007	35375	A	5048	614	1236	TARVAAARRRCQCWGSGACGS ALTLPRTSSLTLTETPIVQQTRM VVRCPTASLPSQWRSCPSSTWA VPKTPPTWTCWRNSAILSL/VVF IRDKLMERRNRRTGRTEKARIW EVTDRTVRTWIGEAVAAAAAD GVTFSVPVTPHTFRHSYAMHM LYAGIPLKVLQSLMGHKSISSTE VYTKVFALDVAARHRVQFAMP ESDAVAMLKQLS
5008	35376	В	5049	1	1281	
5009	35377	Α	5050	2	893	
5010	35378	Α	5051	1	1416	
5011	35379	В	5052	49	768	
5012	35380	Α	5053	1	792	
5013	35381	С	5054	1	1461	
5014	35382	Α	5055	887	1203	RPEKAR/IWGVTDRTVRTWIGR AVAAAAADGVTFSVPVPPHTF RHSYAMHMLYAGIPLKVLQSL MGHKSISSTEVYTKVFALDVAA RHRVQFAMPESDAVAMLKQLS

SEQ 1D	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
5015	35383	A	5056	740	1431	CCQTLPVFHPPHPSHLGPRPCPP
	ł					TH/HLPSEHRSLFLEAACHDSDS
						LEPLNLSSGSKTKSPSLPPKAKK
						PKGLEISA/RPAGALRHRHRLHR
						PQQPSPPLGIPHPSLL\TAQVFIR
1		ŀ				DKLMERRNRRTGRTEKARIWE
						VTDRTVRTWIGEAVAAAAADG
ŀ						VTFSVPVTPHTFRHSYAMHML
						YAGIPLKVLQSLMGHKSISSTE
						VYTKVFALDVAARHRVQFAMP
						ESDAVAMLKQLS
5016	35384	Α	5057	1040	1477	
5017	35385	A	5058	I	1422	
5018	35386	В	5059	1	1404	
5019	35387	Α	5060	353	870	AWSSPRSRCPLHSRTPSLPRVEG
						QAHIGPGL/HRYQQGHQDLFIL
						RSDLPSQVFIRDKLMERRNRRT
						GRTEKARIWEVTDRTVRTWIGE
		1				AVAAAAADGVTFSVPVTPHTF
		1				RHSYAMHMLYAGIPLKVLQSL
						MGHKSISSTEVYTKVFALDVAA
		ł				RHRVQFAMPESDAVAMLKQLS
5020	35388	В	5061	1	1612	
5021	35389	Α	5062	2	429	WGHWASPRLWWAWGSPGPAG
						LKPPGTVAASTGFSQWLLYIETI
		1				SRTPVELELKNSKVRKDVLS/R
						GPHSLYLVQDAQLEHAMEIAK
		l				VSVQALQAYRAPPPRMEQANP
						KTPPDANLYKKGPSPSPRRANT
						TRLGSRATKTIHPDS
5022	35390	Α	5063	508	1007	PPLAQKQAVASQSHSPCGGFHR
						RWFRMRLTGEIMKIITCFKLVP
						EEQDIVVTPEYTLNFDNADAKI
		l				SQFDLNAIEA\ASQLATDDDEIA
1					•	ALTVGG\SLLQNSKVRKDVLSR
1						GPHSLYLVQDAQLEHALPLDT
}						AKALAAAIEKIGFDLLIWTSSRI
						QAVALASPLSNT

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
5023	35391	Α	5064	<u> </u>	1670	 MNIEFFDPPRIVSGLTLNRINIQH
						PFHLTGIRYCIDERORTVNDVL
		ļ				QAHAGRVLQQYLPAPATATFR
		l				CQSRNRTENPQFLVCQWAGED
						QITQLLQEQWRVLIVMIFRTHN
		1				HMAGPRRWFRMRLTGEIMKIIT
		l				CFKLVPEEQDIVVTPEYTLNFD
		1				NADAKISQFDLNAIEAASQLAT
	ļ					DDDEIAA\LTVGGSLLQNSKVR
		1	:			KDVLSRGPHSLYLVQDAQLEH
]				ALPLDTAKALAAALEKIGFDLL
		1	ŀ	!		
						IFVKGTCLVLLVAFGLVTTHFQ
	1					PCLFGVQNGEQETVQILLVGVV
:						GGMERAERIIIPFTLAINNQPTT
	1	ŀ				RLTKTIAQGMPAIIQRNIPNPRIS
ļ	ŀ					LQPAPTPRHSANPATMQPHAQ
1						ATPTPKPHTRTYPTRPPHPQRSA
						TPGTQRTPTPGPQTARDRAPKP
						DHTRPQTAHTAHHTSNDPQQR
						HRTRQRHTNKQHPHPAPPRLPT
						LPQNNKRQKPAESDTRKHHTP
						HMRQPTAHAETTARNTQAAQH
						CHTQNTPQANRHATGDRPEHH
		ŀ				QARDRTRIDPKPNPDPRTITGHT
						KGTTYAHKHTPLATHHMTRPT
						TGHRHGTHH
5024	35392	Α	5065	548	924	GLLRPRFSVNSPHFNTWLLSAN
		ļ				SINWR*YGFMPSLLILLL*LNPM
		ļ				GRYSGASLVSRRMASTTLPGHS
						IAMRQPVCGIFALTRAIISI/AM
						WDFHVEDFMPERMALNLVLNS
			}			AI*LPKIFPARWMKFS
5025	35393	Α	5066	33	424	CRFDESFWCDA\HAIFVAPTLY
		1	•			AYDFYADKTLVPLTDALANTW
						QKQFPLDYGKGRLWGSVAFVI
		1				GSALTGKLVTMFDYRVILALLT
		1				LGVASMLLGFLIRPTIQPQGASL
	į			•		QQESTGWSAWLALVRQNWRF
		1				LA
5026	35394	В	5067	1	265	
5027	35395	A	5068	142	609	CLFRQVSAPHITRVCITRSAATI
		1				QTIDLNNLDTGD*LQPAFPPVW
						LISPGPFHPTPSVTRVPLEKEIRT
1						FAQRLSVLHGINAPEFFDKPLFS
						SLVLTLRDEGYISDSGDAEPAE
						TMKVYQLQTNIVIIHKCFKAGF
ĺ						l ' '
L	<u> </u>	<u> </u>		<u> </u>	L	TLAVGNQFLLINKQRASHQN

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5028	35396	A	5069	983	1387	LVWKVIPWRPAWKITLCVLNS CRRACLFAPAT*TANRLQAVC/ WLQSANPSINRGTLEKESRTVA QRLSVLHGINAPEFFDKAVFSSL VLTLRDEGYISDSGDAEPAETM KVYQLLAELITSDVRLTIESATQ GEG
5029	35397	Α	5070	1	1815	
5030	35398	C	5071	1	354	
5031	35399	Α	5072		678	MTDKQQGEKQKQKTEDKTKT KQKQRKNKTDKRRKTKTERNT QRTPERQTNTNQTRTKQ/RRPD TGVTQADTRQDTSARR\LMRQP RSAAAWMAALRGARDPKLITM VPHSKISRSIVRKGLPMTETQRR TETSKNYRPREQQTRNRQQQD PRQQRTVQTHQQEERRETKDN NTHTKPHETLARRHRRHNIHAL AAADLVIEAASEVWKSKKRSL HSWRKFAPTNAIDH
5032	35400	A	5073	93	729	SADPRNRRDTRAAKFTRDAGK TDC*NL*THIETPDPGDRYSRAG SCGPG/PLKRRLNVWKSKKRSL HSWRKFARHKRY*PLTLRQSLY TAIAAEIKNPERVAGLHFFNPAP VMKLVEVVSGLATAAEVVEQL CELTLSWGKQPVRCHSTPGFIV NRVARPYYSEAWRALEEQVAA PEVIDAALRDGAGFPMGPLELT GPDWAGPPFCCHLFGV
5033	35401	A	5074	145	1038	SILRLRQKRKQLTAYDFADNCD KALY*RNHRLIREILALVCAHD RQKK/YLPSLAGFL*LKNEINRL SVRGW/SRRRSFSVR*/PFTNTD RARSHFYQFVIIDELQSLFQSHADRNQNNGFVSTGSTHVGQFL TGQAVYSQVVRAAMNTDNLTF VNFCTVTEEQLTAILQTEQRER DRFTLTVRDQYAVLTLTHFTRT YVVVVAEGGVQQTSTGSHGHELRTEANQTTAWDHVVETYAAFTVWIHVFQVALTFAQRLHYRTLMLFFNVQSHVFIRLLFTTVDFTEDNFRTGYRQFEFFTTH

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5034	35402	A	5075	39	4329	YAPATFSAATQPTPKKINNPVP RNSAMHWQTRIQAQANGYWL EREQWLAVGEQWFWRMEWSG PHRAAAGRLCHHHGNDATWH RG*FHLCFRPCV**RIEQPQEGY AIIMETTPPGTGVNFIFVSDPAF DKGYAFDFFCLEPMSHAPDDH HRPEGGDLIALAPGESTTSEMS LRVPVANCAVTTASLDSYYGE AMAIGERAPVALLDFAASARL AVGEALTNIAATQIGDIKRIKLS ANWMAAAGHPGEDAGLYEA
5035	35403	A	5076		625	MRASGTDEAVVLVPPIRMTLEQ ALEFIDDDELVEGAAAAFEALE MMGSGRDREI\RYGEGSP\WFDI \VLPCGGGITLTLHKLR\SAQPLL \AVLNRLEQKKPVGLRYDPQAQ SLVCLPTQTRTGWNLNGFEASI THRALRTSRYDGQYYCQGPRT LTIRRTVIMTGSSDPHDTTDSNN DRVLGPSRYDGQYYCQGPRTL TIRRTVIMTGS
5036	35404	A	5077	3	516	ATLSCEVVLENLAASPVVTTLE YTLFDGER\VVHSSAIDHLAIEK LTSASLAFTVEQPQQWSAESPY LYHLVMTLKDANGNVLEVVPQ RVGFRDIKVRDGLFWINNRYV MLHGVNRHDNDHRKRRGVGM DRVEKDLQLMKQHNINSVRNA HYRKRRNKEVRTKKKIESIEEK
5037	35405	Α	5078	3	678	ASYYCTRN\DLELVAVCDSRLS QRQALAEKYGNASVWDDPQA MLLAVKPDVVSVCSPNRFHYE HTLMALEAGCHVI\QAREM/CD TARKLGKVLAYDFHHRFALDT QQLREQVTNGVLGEIYVTTAR ALRRCGVPGWGVFTNKELQGG GPLIDIGIHMLDAA\MYVLGFPR /SVEDSLCGTIELHSGCILWLETS FALNIREHAIMNIRFCGDKAGA SVFPAHIYTATQR
5038	35406	A	5079	2	294	YSFMV/PIFGGRRWTVFSTAILI/I PCVWLGIAVQNPNT/PFGIFIVIA LLCGFAGAN/FASSMGNISFFFP KAKQGA\PVFAF/LGVNGVPQA DGSVMSLAMPH
5039	35407	Α	5080	1	834	
5040	35408	Α	5081	2	135	QYASALVSTAIALLLV\CLAFLY PAAN\SEIHL\GVLSIFWGDPDG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
5041	35409	Α	5082	1	324	MQGGYNIHPLIDALDDAKLAPI
				1		AAKALSHTLLMFDNFYDVEEK
		}		1		AKRGNEYA\QVMQSWAD/AEW
						FLNRPALAEKLTVTVFKVTGET
						NTDDLSPAPDAWSRPDIPLHAL
						GC
5042	35410	Α	5083	1	866	MGLAQAKLLIEAPVNGGRNYN
						GPKVAKFLVGGASLNDVYLHQ
						HDKYRYDRYHLIRNREKTH\IM
						PEWVTQNIMNGRGVNRLDDVN
						RPKSPEIKNDIRLFRQELRDAAY
		1				HFQGDADNDQLSVTPLVGLGK
						SSLLIK\TIFHRMRWAEQNLSICS
						SSFNRPAIFVRNIIQLLREGKKV
		İ				EITVGDKHENDFYIPEDEPFKIIG
						ALPYLYEINLRRFLSRLQYYVN
						TDQLVVRVW\KEDDN\TYHL\K
						GWWGDDKWMVITG\NKLDPR
						AWRLDLENAILIHDPQLELAPQ
						REKELE
5043	35411	Α	5084	1	3483	VMFFFLPGLLMASWATRTPAIR
						DILSVSIAEMGGVLFGLSIGSMS
						GIL/SLGVVSETLWDT*C/HSWS
						RCPAH*SG**Y*/VLALWLTSPL
						LFAVGLGVFGASFGSAEVAINV
			1			EGAAGEREMNKTGLPMMHGF
					İ	YSLGTLAGAGVGMALTAFGVP
						ATVHILLAALVAQIAAWYKAL
						QEQIPDFIPRAPQRQMIADVAK
						TLAGEEGRHLAIEAPTGVGKTL
						SYLIPGIAIAREEQKTLVVSTAN
						VALQDQIYSKDLP
5044	35412	Α	5085	1	848	MLVGKVVVSEGASFRTHGAVD
						TSKAYVSLENSGWTIIADITTTN
						QNTLLNLANLAMSDANVIMMD
						EPVTRSSVTASAENFITLTTNTL
						SGNGNFYMRTDMANHQSDQL
				•		NVTGQATAGDSLTLVTTGGGD
		1				AAFTLGNAGRVFDIGTYEYTLL
		1				DNGNHSWSLAENRAQITPSTTD
						VLNMAAAQPLVFDAELDTVRE
		1				RL\GSVKGVSYDTAMWSSAINT
					1	RNNVTTDAGLRAQAIKTVHHR
						QNRLIHFSLNGSTFNVYRHFRR
						TKTCPKDAETNGKEQGRCQPE
						CQT
5045	35413	В	5086	75	1777	

SEQ ID NO:	SEQ ID NO: of peptide sequence	ł .	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5046	35414	Α	5087	1	860	MKAPGLPADQQFFADLFSGLV
						LNPQLLGRVWFASQPASLPVGS
						LCIDFPRLDIVLRGEYGNLLEAK
						QQRLVEGEMLFIPARAANLPVN
				į		NKPVMLLSLVFAPTWLGLSFY
				İ		DSRTTSLLHPARQIQLP\SLQRG
						EGEAMLS\ALTLFSRSPLEQNIIQ
						PLVLSLLHLCGSVVNMPPGNSQ
						PRGDFLYHSICTWVQDNYAQP
						LTRESVAQFFNITPNHLSKLFAQ
						HGTMRFIEYVRWVRMAKARMI
						LQKYHLSIHEVAQRCGFPDSDY
		l				FCRVFRRQFGLTPGEYSARFQG
5047	35415	С	5088	86	601	
5048	35416	Α	5089	1	870	
5049	35417	Α	5090	305	625	VAPGDRHAFPLAPSGLSPELTLP
	ŀ					QTQCCAQATVQGLEGTRSWSQ
		l				SGTSSLSPWSHTSLRRRRKEEG
						EEEEEEEEEEEEE
	1					EEEEEEEEEEEE/EEEGEEEEE
						EEEEEEEEEEEEEE
						EEEEEEAKTQ
5050	35418	В	5091	1	552	
5051	35419	Α	5092	2810	3117	LTLNSHMAPALIPILAMSAPMT
						PMKN/LSQVHAQGLVIRDLPLIA
						SNFRNTEDLSSYLKRHNIVAIA
1			i			DIDTRKHAACAVPNTLVARGL
						RKHLKRVAAPQHWIAG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	1	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
5052	35420	ĪA	5093	2	1902	LKYPMKRVGKRGEGKFERISW
						DEALDTISDNLRRILKDYGNEA
						VHVLYGTGVDGGNITNSNVPY
						RLMNSCGGFLSRYGSYSTAQIS
						AAMSYMFGANDGNSPDDIANT
			•			KLVVMFGNNPAETRMSGGGVT
						YYVEQARERSNARMIVIDPRYN
						DTAAGREDEWLPIRPGTDGAL
						ACAIAWVLITENMVDQPFLDK
						YCVGYDEKTLPANAPRNAHYK
						AYILGEGPDGIAKTPEWAAKIT
						SIPAEKIIQLAREIGSAKPAYICQ
						GWGPQRHSNGEQTSRAIAMLS
						VLTGNVGINGGNSGVREGATG
į						SAVGEVVFNTSMTGYQEILTD\
						LPI\PSNRYS\SYP\ILAMSAPMTP
		1				MKN/LSQVHAQGLVIRDLPLIAS
		1				\NFRNTEDLSSYLKRHNIVAIA\
						DIDTRKLTRLLREKGAQNGCIIA
1		1				GDNPDAALALEKARAFPGLNG
:						MDLAKEVTTAEAYSWTQGSW
						TLTGGLPEAKKEDELPFHVVAY
į		1				DFGAKRNILRMLVDRGCRLTIV
						PAQTSAEDVLKMNPDGIFLSNG
						PGDPAPCDYAITAIQKFLETDVP
		1				VFGICLGHQLLALASDVEKNVV
						MITAQNHGFAVDEATLPANLR
						VTHKS\LFDRYVTGHSSHR*TG\
						FSFQRHPEASPGPHDA\AP\LFD
5053	35421	Α	5094	8	456	
5054	35422	Α	5095	377	1228	KQARGRLCPAKPQGWSCPRPW
						QPTFC/GQCDLKVRPGVIGDHF
						GALKFDCPTGFRTCLGRITPLFL
						PSSPIWNGCIFPIPVPGGDCGEG
		ł			·	VQIRSLSCMVHSGSISHAAGRV
						EDALCGEMPFQDSILKQLCSVP
		1				CPGDCHLTEWSEWSTCELTCID
]		1				GRSFETVGRQSRSRTFIIQSFEN
						QDSCPQQVLETRPCTGGCSPQA
		1				RPAAIRQCIPACRKPFSYCTQGG
		1				VCGCEKGYTEIMKSNGFLDYC
		1				MKVPGSEDKKADVKNLSGKNR
-						PVNSKIHDIFKGWSLQPLDP
5055	35423	Α	5096	1	519	
		1		 		

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5056	35424	A	5097	I	743	MCPAIDAVLVYVNRKWTNLSD PMPVGQMGTVKYVLMSAPATT ASYIQLDFSPVVQQQKGKTIKL ADAGEALTFP/RGVMPKA/PPQL LVPATFLSAELALLQKSFHVNI QDTMIGELPPQTTKTLARFIPEV AVAYPKSKPLTTQIKIKKPPKV AMKTGKSLLHLHSTLEMFAAR WRRKAPMSLFLLEEHFNLKGQ HSLHENQLQMATSLDRRGN/YT GFITSYLEEAYIPVVNDVLQVG LPLPDFLAMNY
5057	35425	A	5098	2327	3435	KTTTLEDNLGNTIQDIGPGKDF MMKIPKANATKIKIDEWDLIKL KSFCTAKATTKRVNKQDESLRS HYE*WGMLTDCVVMRDPNTK RSRGCGFVTYATVEEVDAATN ARPHKVDGKVVEPKRTVSRED SQRPGAHLTVKKIFVGGIKEDT EGFAFVTFDDHDSVDKIVIPKY HTVNGHNCEVRKALSKQEMAS ASSSQRGRSGRGGGFGGNENFG CGGNFSGHGGFGGSHDGGGYG GSGDGYNGFGNDGGYPGGGPG YSGGSRGYGSGGQGCGNQDSG YGRSGSYDSCNKGGRGGFGSG SGSNFGGGGSYNDFGNYNNQY SNFGPMKGGNF/GGRRSGP*GD GGQYFAKPPNHSGYGGSSSSSSS
5058	35426	A	5099	1	636	MRDPNTKRSRGFGFVTCATVE EVDAAMNARPCKVVGRTVEPK RAVSREDSQRPEDTEELHLRDY FEQYGKIEVIEIMTDQGSGKKR GFAFVTFDNHDSMDKTVIQKY HTVNGHNCEARKALSKQEMAR ASSSQRGRSGSGNFGGGRGGF GGNDNFGRGGNFSGHGGFGGS HGGGGYGG/SGDGYNGFGNDG GGGSYNDFVNYNNQSSHFGPM
5059	35427	A	5100		3622	MTGICYTEDERSYKKNAQPTA ASKKQKETQKFCLRVDGQQKV KLSVLQEKSAQLTVQLKSQKFL GHPTAGRGRSELCLDLPDPPED PVALETRSVGTWVRERDLGMP DGEAALAAKVAVLETQLKKAL QELQAAQARQADPQPQAWPPP DSPVRVDTVRVVEGPREVEVV ASTAAGAPAQRAQSLEPYGTG LRALAMPGRPESPPVFRSQEVV ETMCPVPAAATSNVHMVKKISI TERSCDGAAEMKWEDQNIGD

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
5060	35428	Α	5101	1	782	MRRRLQLDSLLHPPSCFSSTLK
		1				CGKAAAAACDSSDQIQERLQT
		1				GHLPCEASHTICPPNHHSQSTLR
		l				PPLPPP\HNHTLSHHHSSANSLN
						RNSLTNRRSQIHAPAPAPNDLA
						TTPESVQLQDSWVLNSNVPLET
						RFWCCQRFLVFPGLQLQHLTLS
						LPVWSHGALPVSLCIFTWLSSH
İ						KDTSHFGLRVHLLSLPLEAKYS
						FYPGCPVLEPAPNPLKGPWCLW
		l				GVGTFRRYCTTFLFNESEPKDL
						VGEKFKSDRLCREAHLQE
5061	35429	Α	5102	3	1864	
5062	35430	В	5103	1	525	
5063	35431	Α	5104	2	351	RPAVGAEKSNPSKRHRDRL\NA
						E\LDHLASLLPFPPDIISKLDKLS
						VLRLSVSYLRVKSFFQVKRQRR
						EIHGRNCQKKRNQNLKDRGWQ
						WQMPKMWMTTRANEGPCRSG
						GPEHWTPRP
5064	35432	Α	5105	1	375	RPLQKQRPAVGAEKSNPSKRHR
						DRLNAELDHL\ASLLPFPPDIISK
						LDKLSVLRLSVSYLGVKSFFQG
:	}					QGLAVADAEDVDDHTGERRP
ļ						MSFRRPRALDTQALRRTQFGLH
						LLMVNIAGLIATDRL
5065	35433	Α	5106	407	910	NVTPAHYDEQQNFFAQIKGYIR
						CILFPPDQFECLYPYPVHHPCYR
						QSQV*LDNPDYERFPNFQNVVG
						YETVVGPGDVLYIPMYWWHHI
		ł				ESLLNGGITITVNFWYKGAPTP
]				KRIEYPLKAHNKVAIMKNIEKM
		į .				LGEALGNPQEVGPLLNTMIKGP
	_	<u> </u>				IQLACQGSRPPAR
5066	35434	Α	5107	237	908	HEKPGRQRETRSQNHHHHHHH
		l				HHHHQEKIRIQTFTERP*EDTER
	•	l				SDCPQAKKRSPRRNQPC*HLDF
						ELPAARAATQPVNSGARAGRA
1						GATVPRHTATLLLHQRRDYA
						GGSLCDIDRAPDAKQGGKS*ES
						ETDSRFWKGTVC*DHTAGWVH
						GRPTTPHHSSPDSRNGGFQLHQ
						QIALAAGWREGASRPKQEPLQP
						GPEQGQRRGRREPVQGKAPSG
				_		HQSWPAE

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5067	35435	A	5108	57	873	SGLCTAAFHHQMEVVHT*SVSS RS*R\PTTPVIAQWAHEQSGHG GRHGGYAWTQQHGLPLTKAD LAMATAECPICQQQKPTLSPRY GTIPQGDQPATWWQVDYIGPL ASRKGQRYGFAYLARNVSAKT TIHGLMECLIHHHGIPHIIASDQ GTHFVAKEVRQWAHAHGIHW FHHVAYHPEAGGLTEWWNGL LKSQLQRQLGDNTLQDWGKVP QKAMYALIQCPIYGIVSPIARIH RSRNQGVKVEMAPFTITPSDPL
5068	35436	A	5109	1	311	AKCLLPVLVTLRSAGL QITMQPELPIMNWVLSVPSSHK MGHAQQH\QEVAQMPRVSTPA TLPSLPQPALMAPWTVPYDQL MEEEKARARYAGTIQKWTAAA LQPLSRTSLKDSGEGTSKW
5069	35437	A	5110	3	591	DPADPMVLEVSEADRDA\VPIS ESQQRPLGFWSKALPSSANNYS FFKRQLLACYWVLVEIEHLTM GHQVTMRPELPIINCVLSDPCSH KVGHAQQHSIIKWRWYIHDWA EGTSKI.HEEVAQIPMVSTPSLP QPAPMASWEVPYDQLTEEEKT RAWFTDGSARHAGATQKWTA VALQPLSGTSLQDSSEEKSSQW
5070	35438	Α	5111	1	582	
5071	35439	A	5112	1	635	

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
5072	35440	A	5113	2	1538	FFPIGFWRQHIPRLSVLLQHIY*
						VT*KAASFEWGPKQEKAL*QV
						QAVVQVALPLGPYDPADPMLL
			ļ			EVSVADKDAGCSLWQAPIGES
						QWRSLGFRSKALPSSADNYSSF
						ERQLLACYWALVETECLTLGH
						QVTI*PELPIMNWVLSDPSSHKV
						GHEQQHSIIKWRWYICDWAVV
			-			GRESTTNGLAGWSETWKKHH
						WEMGEKQIWGRSMWMELSEW
		•				SKTVKIFVSYVSAHQCVTSTEE
			•			DFNNQWDGMTCSVDTTHPLSL
						TTSVIAKWAHEQSDHGGRYGG
						YAGAQQHGLPLTKADLAMTTA
						ECPICQQRPSLSPQYGTISQGD
				ļ		QPATWWQVDYIGPLPSWKGQR
						FVLTVIDTYSGYGFAYPEHNTS
						AKTTVNGPMECLIHRHGIPHGI
						ASDQGTRFTAKEVWQWAHAH GIHWFYYVPHHPEAAGLIGWW
						NGLLKSQLQCQLGDNTLQGWG
t						KVLQKAVCALNQHPIYGTVSP
						VARIHGSRNQGVEVEVAPFIITS
						SDRLAKFLLPVSSTSCSAGL
5073	35441	A	5114	488	966	ILVGSKGQSVCPGPSSSGTRGSS
13073		ľ.	' ' '	1.00		SPGVAVSNKAAATVGPQ*VTQ
						QEGGPGSRQRTRAGSPELVPAG
						PSAVEGEEGSWTPGGPQVPRKR
						SAAPSRQR*HWLRCTPA*QRRG
						DWPRAAGRCHAARPPAAGSRA
						PTKISARPTRISRLGPRAASKAQ
						RLQRL
5074	35442	A	5115	197	421	KLKATMNSFHKS*SCALVSTRP
						CSPRGDPRATLTTIPALVEPPCG
						TQDPKMAAEAPQRPGKRLGRN
						LLTRRRPT
5075	35443	Α	5116	3	5649	LLEKLRQRIHDKAVALERAIDE
						KFSALEEKEKELRQLRLAVRER
			•			DHDLERLRDVLSSNEATMQSM
			[ESLLRAKGLEVEQLSTTCQNLQ
						WLKEEMETKFSRWQKEQESIIQ
ļ						QLQTSLHDRNKEVEDLSATLLC
						KLGPGQSEIAEELCQRLQRKER
						MLQDLLSDRNKQVLEHEMEIQ
						GLLQSVSTREQESQAAAEKLVQ
						ALMERNSELQALRQYLGGRDS
						LMSQAPISNQQAEVTPTGRLGK
5055	1	<u> </u>	5.15		1.57	QTDQGSMQIPSRDDST
5076	35444	Α	5117	2	157	YLGRHLCLFSEHEEHSSRKEMV
						LVEGLCSEQGRRG*TLASSSER
		<u> </u>		I	<u></u>	KPLENER

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5077	35445	A	5118	248	621	TDEPSLLSHTFCPKGALLSACSG ACPLSSGTQAGMPPSTEALQPC RGLTVPHQ/LG*RPQGMLWWK ALAADWSVEWQKGRRGLGKA LSVTSRTGRLLCR*GSAEAAAS PLAASPRRSSRALGAGIP
5078	35446	A	5119	2	157	YLGRHLCLFSEHEEHSSRKEMV LVEGLCSEQGRRG*TLASSSER KPLENER
5079	35447	В	5120	125	994	
5080	35448	A	5121	1	550	RRYRPRRVQDPGGPAEAERGPP RRLLNARVHVCRAEASERAPGP CGRVSAHVERARDPRLE*QLED KAAPRQDFTQQLRGPEKQHRM VWQEKEDMHKQLVEASETLKS QAKELKDAHQQQKLALQEFLE LNELMAELYSQKQKVWDKEEE MEVAMQKADMMWQEI*RSKK LRKRMLFSQMRWL
5081	35449	A	5122	135	638	GSFPGPAPLGQTLVTLGLAWQS SGWILRGWLVPPRAKCSPRGQ ALAICLSLRGLGARSTWAETRP QGPRGSLASLCPAHMDLSIWKT GDGGPRSAFSKVHQGSLNTFG VDTAARRLSQARTAPSPSV\SAP SSAAAPASAGPPPAPGAPPTPA QALRPAPPSPRPAER
5082	35450	A	5123	193	610	VGELLPFRFKISLRPFLALCLSSL GPCRQPHSGSVPA*NPLLGPRA KSSPHSGCGLAPFAIPELLRRLP LEGSAHCPAPEGAASPPTSWDS ADPPCRETSIFAPPRVVLTPGPS APQASPSQPLLFLTPRWDRLPES VL
5083	35451	A	5124	2446	2630	VGFFFFFWWGERGTGSYSVTQ ARVQ*YDLGSLQLLPPSFKGFS CLSLPGSWYYRHGPPRLG
5084	35452	A	5125	3	710	EDGDWDQRRGGEAGDQDQRR GGEDEDQDQRRGGEDGNRDQ KRDGENGDWDWKREDENKDQ DEKREGEDGNQDQRRGGEDGD QDQRRGGEDGNQDQRRGGKD GDQDQGRGGEDGDQDQGREG EDGDQEQRRRGEDGE*DQRRG GEDGGQDQKRGGEDGNQDWS REGEDRD*HWRRGGEVGE*DQ RRGGEDGE*GQRRGSEDGDWD RSRGGEDGDQDQRSVGHLCHS QGPPASGCVS

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5085	35453	Α	5126	506	882	RCSGQGGFDGNKGTKALPTISA
						DTTRCWRTLRVTQMTCRFRRF
						KAFFIGIMSCGRTGRILFPPCSS
						MSWIPCRAKNS*GNVISQSPSK
						NR/WQIMMVIQLLNFYLPGDFV
						SFRIMLKGNWEVTRTRG
5086	35454	Α	5127	107	1327	ERRKRAVGGFDLVLGEQPSDKI
		ŀ				FR*VIV*GLALWLPLSKSFVIPP
ļ						AELAINPSAKCKTDMTVMEDA
		İ				VEVR*VWLQD*GLDSLEVLLDS
						FGPVRDCSKDNGGCSKNFRCIS
						DRKLDSTGCVVGLPCKLVVDS
			•			SGCYDRHIGVDCSDGFNGGCE
						QLCLQQMAPFPDDPTLYNILMF
						CG*VELPARPLDGRSCQLITETC
						PEGSDCGESRELPMNQTLFGEM
}						FFGYNNHSKEVAAGQVLKGTF
						R*TSYT*N*YQQLPDGLVVATV
						PLENQCLEEISEPTPDPDFLTGE
						CAFSTLFGYPVLQHWKVRSVM
		·				YHIKLNQVAISQGEHCCRLSCD
						GATSRADFVALLDQFGNHYIQE
						AIYGFEESCSIWYPNKQVQRRL
						WLEYEDISKGEWHACWQILSTF
						PWIVISLHNCLP
5087	35455	Α	5128	3	23	FYSAFLVADKVIVTSKHNNDTQ
						HIWESDSNEFSVIADPRGNTLG
		1				RGTTIT*VSIPPSL

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5088	35456	A	5129		2462	MRALWVLGLCCVLLTFGSVRA DDEVDVDGTVEEDLGKSREGS RTDDEVVQREEEAIQLDGLNAS QIRELREKSEKFAFQAEVNRM MKLIIHSLL*NKGIFLERLISNSS C\ALDRLSLISLTDENALSVNE* LTVN\IKCDKEKNLLHVQHTGV GMT\REELVKN\LGTIAQIWDQ ASFLNKMP*SHRKMGPVNLPEL IGPVWVSGFLFPPSLVADKVIVT SKHKQPIPQHIWGVWTPNGIFL *LLDPRGKHS*DRGNRQFTLCP *KEEASDLPWNLDTIKNLRQKN IHSFIKLFLIYVW\SSKTETV*GA PWGEEEAAKEEKEESDDEAA\V EEEEEEKKPKTKKVEKTVWDW ELMNDIKPIWQRPSKEVEEDEY KAFYKSFSKESDDPMAYIHFTA EGEVTFKSILFVPTSAPRGLFDE YGSKKSDYIKLYVRRVFITDDF HDMMPKYLNFVKGVVDSDDL PLNVSRETLQQHKLLKVIRKKL VRKTLDMIKKIADDKYNDTFW KEFGTNIKLGVIEDHSNRTRLA KLLRFQSSHHPTDITSLDQYVE RMKEKQDKIYFMAGSSRKEAE SSPFVERLLKKGYEVIYLTEPVD EYCIQALPEFDGKRFQNVAKEG VKFDESEKTKESREAVEKEFEP LLNWMKDKALKDKIEKAVVSQ RLTESPCALVASQYGWSGNME
5089	35457	A	5130	45	416	RIMKAQAYQTGKDISTNYYAS RWGLAMFPSLDNKSETLSQKK KKKDLCPHISFEMDLTTADAR WVQDAATRLLATTRCPPNSRG SR**ISWRAGCCPGGWAGPHW GHDGCL*PSCVSSS
5090	35458	A	5131	3	477	GRLPRHRGGEPGAAQPAGRQP RGGAAATAAGAPRAGGGAAA GNARRCREAAARDPHHGEGSG PEPSQCEGAAHQRAGRAQGD* GGSGATGEGGRRGHDSHNPLG RVRGSTLPPS**N*VGL*VPSWP QCGPAHPPGQHPALQEIHQRLP LESGGHRVV

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5091	35459	A	5132	3	446	RTGSWGWSASQQRHRKDRGL DSAGSGWVSHILPVPAPPPADA RPSASRWRRQGSHPLPGRTECP SVQPLCLPPPAGHHRNHSIASPS QTPPPRSLPGSQDKSLFLPRDRR TGEGEGHMN*APLSAPGTSCRS
5092	35460	A	5133	352	588	GGSTSWLPEILGPTQ DIFKVGGCFLFKRLRCQQRFFL LIFSSSTQSA*RVSNGPYH*EIK MAVYYGELRKNGLIQESCDNIQ PRQGSFNMDSMQ
5093	35461	С	5134	13	144	
5094	35462	A	5135	587	1078	LSQRFPLSSRIKLQRLSLQNGIP QIKELDYLFRTNSNQINFNLLSH RVRARPPAPPGRPNPSPGPDRPP EPPARAGSSRRGPGEPPRRQPQ RRFRRRGPDRPGRLSRGL*RRR ARPASPQPRQHHSRGPAGRPGV APRRPAGLAAAAERLTARGAA ASPAIPRA
5095	35463	A	5136	44	455	HLRNRTR/PSQITPHIYNHLIFDK /PLFNIWWWENWLAICRKLKLD PFLTPYTKINSRWIKD*NIRPKSI KNLEENLGNTIQHLG/IGKDFM TKTPKAMATKAKIDIWDLIKLK SFCTAKETIIRVNRLPT*WEKIF AIYP
5096	35464	A	5137	101	861	LIGYQPKMSRTRRIHSRILPERIK YLGSQLIRDVKDLLKENYKPLL NEIKEDTNKWKNIPCSWIGRISI VKMAIL/PQAICGKLKLDPFLTP YTKINSRWIKDLNVGPTTIKTLE ENLGNTIQAIGMGEDFMTKTPK AMATKAKIDKWDLIKLKSFCT AKETTIRVNRQPTEWENIFTIYP SDKGLISRIYKELNQIYKKK\TR NPIKK*VRDMNRHFSKEDIYAA HRHMKKCSSLLDIREMQIKTTM RYHLTPI
5097	35465	A	5138	3	314	LKSFCMAKETTIRVNR*PTEWE NIFAIYPSEKGLIYRIYNEATNK DLVRRSGPLKTPKKKSQLTVLN LVTFKRTPALSEEKESAQVLWQ FKKPECVLTSKRVH

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	I .	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
5098	35466	A	5139	1	1278	I MNIDAKILNKILANRIQQHIKKL
						IHHDQVGFIPEMQGWVNICKSI
						NIIQHINRTKDKNHMIISIDAEK
				1		AFDKIQQRFMLKTLNKLGIDGT
						YLKIIRAIYDKPTANIILNGQKL
						EAFPLKTGTRQGCPLSPLLFNIV
				•		LEVLARAIMQEKEIKGIQLGKE
						EVKLSLFADDMIVYLENPIVSA
						RNLLKLISNFSKVSGYKINVQK
						SQTFLHTNNRQRESQIMSELPFT
						IASKRIKYLEIQLTRDVKDLFKE
						NYKPLLN
5099	35467	Α	5140	1	430	GGVFFLRTPTDCCGMLQGAAG
		-				STGFYASEARPRGPACSLDRGS
						SSLLSWGLHQDPLFPAAKQHRR
						EAGRADQNSPRPRLSTG*QTPT
						L*RRPAAATGRRRAQPPRATSV
1		İ				SAPASPPCPRARPPHHARPPRVL
						ACQP*ATAPGP
5100	35468	Α	5141	1	346	
5101	35469	Α	5142	3	1336	
5102	35470	Α	5143	5	283	
5103	35471	Α	5144	1	1368	
5104	35472	В	5145	l	838	
5105	35473	A	5146	11	962	
5106	35474	Α	5147	19	87	
5107	35475	Α	5148	41	831	GLQNPKHQSSQLVDLKMKKSL
						LQCMFQYLIVLQ*YLK*YLGTQ
		l				LTRDVKDLFKENYKPLLKEIKE
		1				DTNKWKNIPCSWIGRINIVKMA
						ILPKDPTTAQLSPRGSTAQKGH
						WRPCPLPPRGVPRSPRGPTSHP
						HHDPCPGGAPVRNTGSERWQR
						KLRSGPKSDWAGRAQAPSLGE
						GGAKNGKSHPGSHRAFSLPRAP
						RRLGPGSQPGRFRGFLKQARGR
						AGEGHSAILLAPESPNAQVSNV
	<u> </u>	<u>L</u>	<u> </u>			TSATYRSNLSGLPRPCSVVLLG

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
5100	35476		5149	1	12233	I MKAEIKTFFETNENKDTTYONL
5108	334/6	Α	13149	'	2233	1
						WDAFKAVCRGKFTALNAHKR
	İ				1	KQERSKIDTLTSQLKELEKQEQ
ŀ]			THSKASRRQEITKIRAELKEIET
						QKALQKINESRSWCFEKIHKID
ļ		1				RPLARLIKKKREKNQIDAIKND
						KGDITTDPTEIQTTIREYYKHLY
		1				ANKLENLEEMDKFLDTYTLPRL
		1				NQEEVESLNRPITGSEIVAIINSL
		l				PTKNSPGPDGFTAEFYQRYKEE
		l				LRIKYLGIQLTRDVKDL\FKENY
	1				ļ	KPLLNEIK\EDTNKWKNIPCSW
						VGRINIVKMAILAQGNL*IYKFN
	1					AISIKLPFTFFIGL/EKNLTFKFIW
	ł					\NQKRS\RIAKSILSQKNKAG\GI
						TLPDFQTITYKATVTKTA\WYW
	1					YQNRDIDQWNRTEPSEITPHIYT
		l			,	ILI\F\DKPEKNKHGGKASL\FIK
		l				W\CW\ENWLA\ICRKLKLDPFLT
						PYTTINSRWIKDLNLRCQIIKTL
						EENLDNTIQDTDIGKDFMTKTP
						KAMATKAKIDKWDLIKLKSFC
						TAKETTIRVNRQPIEREKIFAIYP
ļ						SDKGLISKIYKELKQIYKKKNK
		İ				QPHQKVGEGYEQILLKRRHLCS
	}					QQTHESMAHHHWPCREMQINT
	1					TMRYHLTPVRMAIIKKSGNNRC
						WRGCGEIGTLLHCWWDCKLV
		1				QPLWKSVWRFLRDLELEIPFDP
						AIPLLGIYPNDYKSCCYKDTCT
	Į.	1				RMFIAALFTIAKTWNQPKCPTM
						IDWIKKMWHIYTMEYYAAIKID
5100	25477		5150	1200	2401	DLLTRTACAVPGWGCSSRATC
5109	35477	Α	3130	1390	3491	QARGWRLLSSPPHPSGVQPLLC
						HLPYPPSPFSLLSNLTPFLRAFPR
	1					!
	1	1				SLETS*KWART/WPGAVAHAC
		l				NPSSLGGRGRWIT*GREFETNL
		<u> </u>				TNMEKPCLY
5110	35478	Α	5151	310	445	YLTKIWPGAVAHTCNPSTLGGR
						GGWIT*GQEFETSLANMVKPHL Y
5111	35479	A	5152	44	1132	POVIHPPRAPKLLELQEPPRWIH
						DLNQDTPSALCQH*RTC*HRAE
						GVSWFKSWIHLGGSCNSSSLGA
						RGGWIT*GQEFETSLTNMAKPH
5112	35480	С	5153	183	323	ALGO WIT GOLD I SOLITION IN INTERNAL IN
5113	35481	В	5154	504	2474	
5114	35482	С	5155	66	269	
5115	35483	В	5156	2414	3096	
2117	133403	<u> </u>	12120	2117	12070	L

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5116	35484	A	5157	9	239	HHHYCQHHHHHHHHHHHH LPAPCRNLRV*PVCFRYKLWFH YGKQHLLFLDFLLSHSEWCEM VSHCGFDLHFSDD
5117	35485	A	5158	3	387	TDTKPLSCSSCPPRGKKPVSHR DQ*RRWGRRCRGEPGTEACWS AGALQF*KPRSRGGRTSKMTSG PSAEKGGG*AGRKPGRRHFGRR VLRHSCVLPVARELWGRHGPC GWRQKNVPWKWSELLPVVAP
5118	35486	A	5159	1	277	MPPPPPPSESPGPGPVGSRLPPG PAVTGNEKELRGNP*VGLAPPR PRFRASRREGVTGKVHRPNRLQ LRAPPKG*SDQRGQNVPPSEAK SLC
5119	35487	Α	5160	1	3285	
5120	35488	A	5161	3	194	IPLSLQSKEVGQQLQDDLMKVL NELYSVNQMGCLAL*QAEGSSI WQLAAIVHQQQN*TPSWGQ
5121	35489	A	5162	520	844	VQAPGKCHAQMPAPAAPVPLG RSRNPGRLASAVPVSSLLRFLPF LQPIPPSDILPLSFEAVPSSPAAE SGLHPCTRRRKRARPSC*VPGL LLSPQPHPPPHVWRKGIL
5122	35490	A	5163	1	794	FRGFLDRGDCAALPCTYPHSPC SH*GGNCLPSLLTRPCVKA*PQ MSGRKSSMRRWRRQSRLTAGT SS*TPTSSTMC*ALVGSSTWNC MLQAGSTAPGAGTPGSRPTWS SSSTCSWTAPSGRARCACASSS SCAMSAARRGWTSPACWRRTS RAWWTTSSPACASSATASVAA STASTWPAARTTGGTAESSARP ARRASCTGSPARSCWRRRPPT PSPGRPAPPSRRTRRAQAGTSA LSPGACFGPRSCC*SSTCSSLSV APY
5123	35491	A	5164	3	57	RIGQGVPVVHS*VEGGPNVISIV LEYLRDTPPVPVVVCDGSGRAS DILAFGHKYSEEGG*VKVFLWC THKWKEDPM
5124	35492	Α	5165	1	1326	
5125	35493	A	5166	2	337	YSVTMTPGKLRTLCEIDWPALE VGWSSEGSLDRSLV*KVLCPEL VPSGGFLVSLTSRMKPQTLTGS VKVLKDGMSGVCSFRCSDVSR VSSFWWVRGLAGWSTTLLRKL
5126	35494	С	5167	339	548	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	İ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	}			sequence		
5127	35495	Α	5168	3	413	SQAFLPAASRRERPPEPVDPKA
						TDPRSGTRRLPPRLPCQWTTPE
						HSHTSQLNLSRAFPKEPNPAAT
					•	Q*NPSGASPRERPHARAQEDRN
		ŀ				GEC*EGNGSRPDLEEEASTPRA
		1				SPTPSD*PLWLHPPRPLVLETPL
				•		LQL
5128	35496	Α	5169	3	479	FLHSTHIYWAHSMHELLKTAV
						VSISGQCTSLNTESGIWGGVPE
						NAR/CPPAEGTVPASLQPRDPPR
						LPGSH/ASLPSASPPSRHVTWPP
						EP/EAAGPKPS*IRCRWPPAGAT
						TKANPKESGRLVESMLPLPKSR
						SAPVTFSAFACSGLPERVHEVF
						RVGKLR
5129	35497	Α	5170	81	329	QSVSSPSWFPQFPVQVQKVPCY
		İ			<u> </u>	QVYRGDLVV*KHVAPPPSLPSS
		l				FSGHVRHDCFPFAFHDDRKFPE
						ASPAMLPVSSLGPHYL
5130	35498	Α	5171	335	467	MPSTPTARKPESPEALARVESIA
						CLPSPAP\PHECSPQASEETG
5131	35499	Α	5172	1	2424	
5132	35500	Α	5173	2	304	CLTKNTSATREKQVTELHRRSK
						LQPEPAKTKALQTVIE*KDITLC
						*VKEHLYIVACWALRLHLVFLP
						TKAGISPALMKVSYPGKPLTLK
						QLGRLVILQEQ
5133	35501	Α	5174	44	286	GVQNGVEELILVRRMQKSPGP
						GEMESGSLEKEPLGTQTGPVPS
			Ì			EGTE*TQGPPRTPSLQTQGPQRT
	1					PSLQTQGPPNTPSL*TQGPLRTL
						SLQTQGPPNTPSLQTQGPPRTPS
						LQTQGPQRTPSLQTQGPPNTPS
5134	35502	Α	5175	206	384	KRVFAHQADGRARSGFISSAGP
						GSQAGPGQGTG*QREDVEAAG
			<u> </u>			QPGSTAEGVRT*SDPG
5135	35503	Α	5176	270	818	RNERKPSQRTAVGLGAGQLQP
						GQPGVQSEEPGPLEAGCAAGV
		Į.				LPGPAEQDREGIYSK*KRVFAH
	Ì	ľ				QADGRARSGFISSAGPGSQAGP
						GQGTG*QREDVEAAGQPGSTA
						EGVRT*SDPGSSHPAAAAFYEN
						IQDGEALGRCGAKRTATAAP*Q
						RG*EGF*AGPTAAEKNEEGPPP
			<u></u>			VFSLFNSHL
5136	35504	Α	5177	122	436	LTSLFSAISTKAKVSPLCSARAC
						SPSPAARAALGSPGFTC*GAGV
				1		APTPVPVAAPAPATAPATAVAA
1						AATTTVVSAAMFTRRATRAAA
						ATGAWGFVAGAPWPLQT

SEQ ID	SEQ ID NO:	Met	SEQ ID NO	: Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence	İ	
5137	35505	A	5178	20	1341	MRRCVAAPLAHRW*QKLGLPG
		1.	3170			CTPTVCSFLSPQPPPRARSSSAT
				!		LSSGAPRRAATPQPQTTPPSSVQ
		ŀ				PCAATPCARGGRPAPAGVTWP
						TTRPSMA*RTS*ASTTAPRMAP
		1				PRSHACARSHRPETARSARTAP
						RSAITRRAFTSTRPPPTTRTVAS
						· ·
						SGTHTSGLSPTASRLARCRAPG
						RSSTIIT*TCRSPTRLCCPAQRPL
İ		}				PPASSPSSSRTSRSVWTRRCTRL
						RWTSSRPPSWMALRTVGTSTGP
						TA*RSLRRCQASTWRSRPSTSA
į						PPSWCARWAAT*PLPSACQRK
						WSMLWRTGTARVSTSACGAAP
						STSRSTSRPSTPMLRAPVPAGW
						QPPALHPQPPRPSHTRQPWPSA
						RRSCRWRTCTTRPASSTSSPRA
						T*TSHWPPTTRWRMSRCSTPTK
			:			TNCTCMRGLGTCQAGRLRGCP
						WPPGPSWAPWSRSWPCSLCSA
5138	35506	A	5179	47	598	 ASGPASIVVLRQQFIKFFLDEICS
15.50	32200			,,		LEERVWCRKLCLLYILWKEPR
						AVQAKSRCSDIRRELYRPVSLA
						GLSGDFLRAGRYSAHTVNGAW
						SAWTSWSQWA**SRGLLLALH
						NMLKIIIAFCSLQVATRIQSTFIS
						SWHLSPSLLGSRSFSLYMSFGS
						VLCPLFISHDTSSYSTKSFQSFSC
						LGVW
5139	35507	A	5180	236	690	LQNWSYLAEQCWNGGFIYLIM
	33307	 '`	3100	250		LRRFKHKAHSTYNGNSSNSSEP
						GETPTL*LGDRTAKKGKRTRKF
						GVISRPPANKAPEESKGSAGCE
						VSSDPSTELENGPDPELGNGHV
		İ				FQLENGPDSLKEVAGPHLERSE
						VDRGTEHRIPKTDAPLTTSN
5140	35508	A	5181	2	480	WPFWAMVQRPSSAILCGPATLP
3140	33300	^	3101		100	PNCRGACCARCRATGALSAYP
						AIHRPATRPPAPPAPGPLTSPPPL
						SVQNSNPDPLPSARPSPGPAAPR
						_
						HARHSPSELPALTLSARGSSAL
						CTKLKVEAAA*SCSASAWNWR
				-		LLPRNHVDSRCQPEPDGARARD GRGP
5141	35509	Α	5182	3	153	O.COI
5142	35510	Α	5183	3	239	
5143	35511	Α	5184	1	394	
5144	35512	Α	5185	1	435	
5145	35513	В	5186	84	2068	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
5146	35514	Α	5187	149	832	DAGLSWREATRPGAPLHRPGC
						QDASTQMTRPGDPGSSALFPLP
					İ	SP*PEREELPMSSRRSPKPELPG
					•	EACSRRPPTRPRPRTATRGPPLN
		·				GRLPRARLPVSAGAGALGGGR
						CRRRVAPRRVTPRLATRPSGVA
				-		PRGSG*AATTKLSLAFGVTPLA
						LTPWSRTPFRKREERPPF*YPRP
						ATPEASPAHVENFRSNQTRYTS
						AE*SGQQLERPSIAEISAQATVE
				•		PRSTI
5147	35515	Α	5188	45	312	AEAQGGAEAEGHGGGGGPAPS
3147	33313	^	3100	43	312	QGCPARAAGARGWQPAPSPQK
						P\TPSDTTPPP\RTQPPVTQPCHP
İ						VTQPPVTQPPVTQPRHPMKIPPP
5149	25516	_	5100	15	549	DDILLEDES OAD CDD ADOA AAV
5148	35516	Α	5189	15	349	PRHKEPESQAPGRRAPQAAAY
						QQAASPAR*QGGSCSRCPPG*A
						VEAAPAP*GQALGCVSQGPA*V
						GPTRSPHPTAGLSRPHGPPSVRS
						TGWSPSSRPLQAQEAPWPVAPT
						HIPPPLDPGQPHSCGLGAGCQP
						RAPAARAGHP*EGAGPPPPPWP
		ĺ				SASAPPCASAQPVLWLRFPRET
						KT
5149	35517	Α	5190	1	2445	
5150	35518	Α	5191	1	708	FSEGRNRFLNVGPTRFSTSRQTL
						MWIPDSFFSSLLSGRISTLRDET
						GAIFIDRDPAAFAPILNFLRTKE
		1				LDLRGVSINVLRHEAEFYGITPL
						EKIKMEYCFSVCMYS*QVCFE
						MFLVRRLLLCEELERSSCGSVL
						FHGYLPPPGIPSRKINNTVRSAD
						SRNGLNSTEGEARGNGTQPVLS
		ĺ				GTGEETVRLANTIDSKYVAND
						DWLIFQYVDEEDLMNAIKDFSS
						VTKERTTFTDTHL
5151	35519	A	5192	31	330	YGGTATVQRWAESRHVPFPPS
-						QEQNANKICSVCQEEKDSADSY
						GEIPWWQGPERCWPVRLMLAT
						LGDYKWVLTGIGTDS*LGFTIPT
				1		
						DHANAFSAKEIPQ

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	,	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
5152	35520	Α	5193	1	734	MISEVIVIQMLFLHAFSKVFSAL
						IKNNKKPLQDAQIKWKAISGLS
						GGMRRSECNRERDQVTRTQCR
		1			:	DISQK\RGPGD*GMATSKNGAM
						A*QLGPGCQGDASTGNAEFGES
						AHAWSGGGRVPGVTGPDREEV
		l				GRSWQGTSCGGGTSGCGRLQ*
						DGVASRGRW/DPARPREQRAG
						VRQLAAPLGGP*SRRPWEPPGG
						SSCGRYLFAISSWSGVTVAQEG
						QRKRGSGK*RTERSGPGLGRRL
						PPLLLTV
5153	35521	Α	5194	3	449	EGQRSSSPATALSSSGPPSLKM
		l				ATSKNGAMA*QLGPGCQGDAS
						TGNAEFGESAHAWSGGGRVPG
						VTGPDREEVGRSWQGTSCGGG
						TSGCGRLQ*DGVASRGRWDRR
						DRASSGRESGSWLHHSGARSLG
						APGSRPVEAPAAGSEAKTPG
5154	35522	Α	5195	606	841	CTVGFALPVFY*GFSH*CSSGIL
	İ					V*NSRFLLCLCQALVSG*CWPH
	}	1				KMS*GGFPLLLLTGIVSEGMVP
		1				ASLCISGRIQL
5155	35523	Α	5196	620	930	IPFISFSCLIALARTSNTMLNRSG
						ERGHPCLVPVFKGNASSFCPFS
					İ	MILAVGLS*IALIILKYVPSIPNL
						LRVFSMKGC*ILSKAFSASIEIIM
						WFLSLALFI
5156	35524	C	5197	1	819	
5157	35525	Α	5198	591	815	CENTION ADVINUE DONIEL OVO
5158	35526	Α	5199	187	387	FFKTICFMRVRVKRPPNRLCVS
						NMAVYFTWVQAGLHCESYPEH
5150	25.527		5200		1107	L*WSCRLPRRLGGVRL*PLEGE
5159 5160	35527 35528	A A	5200	119	1197 375	ILDISPLSDE*IAKIFSHSVGCLFT
3100	33328	A	3201	119	1373	LMVVSLAVQKLFSLIISHLSILA
						FVAIAFGVLDVK/CLVHAYVLN
						GIA*VFF*GFYGFRSNI
5161	35529	В	5202	1	1053	UIA VEF GETUERSNI
5162	35530	A	5203	2	157	
5163	35531	A	5204	12	176	
5164	35532	A	5205	2	223	QTVLEDRSPRARWIF*G**GTIC
		-				SIPIPSFWWFSSNLSHSFVCRSPP
						SSSPGGLPMCTCVSKFPWAVW
						PFSRY
5165	35533	A	5206	61	308	
				•	*····	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
5166	35534	A	5207	487	898	DFAEVAYQLGISSGVCRTTDFH
						EPRMLLSDRSSGSFVTII\SVFLS
						FSSLKCL*SPLVHDALHQAFHP
}						GKSRLRFLGSMACCMSKFTCPP
						TLSTVPSSTPFSFSCSLIKTRSAT
						LLKLWSISLRFRLHSWSSAFRV
5167	35535	A	5208	9	191	IPQHIYPPKTKPGRI*ISE*TNNR
						L*NCGNDQ*LINQKESRTRWIH
						SRILPKTNREPNHE
5168	35536	A	5209	1	1989	-
5169	35537	В	5210	1	1488	
5170	35538	Α	5211	929	1209	IPFISFSCPIALARTSNTMLNRIC
		1				ERGHPCLVPVFKGNASSFCPFS
		:				MILAVGLS*IALIILRYVPSILNL
						FRVFSMKGC*TLSKAFSVSIEII
5171	35539	A	5212	242	642	FSLKRSFTSLVSWIPRYFILFEAI
						VNGSSLMIWLSVCLLLMYRNA
						CDFFPLILYPETLLKLLISSSSLY
						LW*NSAVNLSGPGLFLVGRLLII
						ASISEPAIGLFRDSPSSWFSLGR
						VYVSRNLSISSRFSSLFV
5172	35540	A	5213	747	885	
5173	35541	В	5214	115	1267	
5174	35542	Ā	5215	209	322	
5175	35543	В	5216	1	3321	
5176	35544	A	5217	390	655	CLFPVHYPRLHAMMHQMKNP
						VLVLGIGGFILGILGVIGGPVSLF
						KGS*KLLGLIETGKISLDHPVPG
						GTPTKMALYEHFMNTKEFGIE
5177	35545	A	5218	3	221	
5178	35546	A	5219	798	914	FVFLWDRW*YPLYHFLLCLFDS
						SLFFSLLVLLAVYQFC
5179	35547	Α	5220	12	460	YLCMLHEVLVLCFSVPSGHFSF
			ļ			SLN*LF*LAVPVTFYQG\LSFLA
						QA*ASQVDFRQLCWQQEFQAS
						GS*LAGLHGSGIH*ARPLGSWL
						QLPFQESEWFCLSGVPGATVVS
						/CTF**SPF*LV*DGISLWF*FAFL
						*WPVMMSIFFMCLLAA
5180	35548	A	5221	3	508	AERNGSRQGVGAGAGDHEVLQ
	33346	` `	'			LLPDKWKQKILCPVQLLRGAW
						SNYYSPDANRQP*VQGNAHLF
						AECWPHGEDSQPTVPTSQGRM
						LAGPARHLPHPHVPSPSGSLRG
						SDSEDTH*GTVAGSGCLHCLAP
						GHLHPGWPPSRVLPIPDRSVHQ
						GPGRKAPPTPPRVTPLFT
L		1	L	<u> </u>	<u> </u>	UFURNAFFIFERVIFLFI

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
1	sequence	1	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
5181	35549	ĪA	5222	 584	1052	LRKAAASGSFVFVAPDPNTDTG
	33317	``			1032	SSQCSESESGSRGCSMGHQRSL
						ОННУННИННИННОННИННИН
	1					HLDTGWRGE*SRHSAGLCPPSQ
						RRLHSCQTSPSCSPPSGHRCI*LT
			İ			GIEAA*CHSL*PYQKDACHHFE
						1
C 1 0 2	25550	 	5000	1614	2200	EHSPPP*RPPHLPRHHLHPQPSS
5182	35550	Α	5223	1614	2308	LRKAAASGSFVFVAPDPNTDTG
						SSQCSESESGSRGCSMGHQRSL
1		1				ОННУННИННИННОННИННИН
		1				HLDTGWRGE*SRHSAGLCPPSQ
						RRLHSCQTSPSCSPPSGHRCI*LT
	1	1				GIEAA*/WS*PLTLSKRCVSPLR
						RAQSSSVTSSTSAQASSAPPEKK
						*L*PERPYSSPHPASVADLQMC
ŀ						LPLNRITGQTNRPLHRAILLRW
į						MRSCNLLTCTVLSNVSFSFSELS
		<u> </u>				YSRNTMCRG
5183	35551	Α	5224	3	407	SVGLILPSSLQHHYHHHHHHH
			1			QHHHHHHHHLHQRKLAPPLC*
		1				ADLPTCAPHQGIRPFGGFPQTS
1						GSPGLPVYAEEQGTERLSLQGT
		}				GCGCQAGLLCWGPPVASTSVG
						FSLESSVSPEQVPLSPGPAGTDA
5184	35552	Α	5225	1	1287	
5185	35553	Α	5226	3	1315	STMALLHSGRVLPGIAAAFHPG
						LAAAASARASSWWTHVEMGPP
						DPILGVTEAFKRDTNSKKMNLG
						VGAYRDDNGKPYVLPSVRKAE
						AQIAAKNLDKEYLPIGGLAEFC
		1				KASAELALGENSEVLKSGRFVT
					=	VQTISGTGALRIGASFLQRFFKF
		Ì				SRDVFLPKPTWGNHTPIFRDAG
		İ				MQLQGYRYYDPKTCGFDFTGA
						VEDISKIPEQSVLLLHACAHNPT
						GVDPRPEQWKEIATVVKKRNL
						FAFFDMAYQGFASGDGDKDA
		Ì				WAVRHFIEQGINVCLCQSYAK
						NMGLYGERVGAFTMVCKDAD
		1		1		EAKRVESQLKILIRPMYSNPPLN
		1				GARIAAAILNTPDLRKQWLQEV
		1	1			KGMADRI\IGHAGLQLVSNLQG
				1		REG\STH\NWATHSPTKLGMFC
İ						FHRG*KLEQVERLIKEFSIYMTK
						DGRISVAGVTSSNVGYLAHAIH
5186	35554	В	5227	57	386	
5187	35555	Α	5228	1485	1883	
5188	35556	В	5229	117	641	
5189	35557	С	5230	i i	552	
5190	35558	В	5231	250	378	
2.70	155550	1	15.55	1	1	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
5191	35559	IA	5232	3	1117	TPDPGQGCSEGGIVHETRPSPQP
2171	33337		3232			LCQGSCAVGCSPLGLLR*RMM
						ALGKIPPGHRGPAFLPAVESAH
						VVPGLPGLTRETLPAPLGPHRPP
						QPPQGPLSTAPPAPLVSPLGPHV
						PGLPGPPPGCPSPLPLRPCPPPS
						WLRPHRPVQPWWGT*GSGWRP
		Ì				RCRPGAAGRNPGLPRACCRGL
						CRGPH*RPGGPRPGHSGPRTSS
		l				WGRTMGNRPRPSASAG**PRHR
						R*TSAERPGSPPQQSASRTGGRS
						PLSHPGSWWGVMGPPGLWAG
						APSMWG*GPGSRVAGGSGRRP
						TSPRPPHYPPHPPGTWPSGAQS
						GRR*QAQTPGAATEG*GGWGL
						PPKPWCPQDQFLGVQDWVLPP
		1				NGARVGGTWAWKRESVLVPG
5192	35560	c	5233	229	522	Norther GGT WHEN THE GGT EVILO
5193	35561	A	5234	1	972	MTAEYYKLKQVVTPIAAAVPD
3173	33301	1,	3231), 2	VVSLLEQINISLCTWYAVIDLG
1						NACFYIPVHKAHQKQFAISRHG
			·			QQYTFTVLLQ/RRDLDLFSLPQ
						DITL\VHYVDGIMLIGSSEQEVA
						STLDLLVRHLHARGWEINLTKI
						QGLATSVKFLGVQWYGACRDI
						PSKVKDKLLHLAPPTTKKEAQH
						LAASFEWGPEQKGLQQVQVAV
						QAALPLGLYDPADPMVLEVSV
						ADRDAVWSLWQAPIGESQWRP
						LGFQNKALPSADNYSPFERQLL
				•		ACYWALVETEATPVITQRAHE
						QSGHGGRDGGYAWAQQHGLP
		1				LNKADLAVATAKCPISQQQRPI
5194	35562	Α	5235	3	246	
5195	35563	Α	5236	3	841	
5196	35564	A	5237	1	746	RDLDLFSLPQDITL\VHYVDGIM
						LIGSSEQEVASTLDLLVRHLHA
						RGWEINLTKIQGLATSVKFLGV
						QWYGACRDIPSKVKDKLLHLA
						PPTTKKEAQHLAASFEWGPEQ
						KGLQQVQVAVQAALPLGLYDP
ŀ	1					ADPMVLEVSVADRDAVWSLW
						QAPIGESQWRPLGFQNKALPSA
						DNYSPFERQLLACYWALVETE
						ATPVITQRAHEQSGHGGRDGG
I.				ŀ		YAWAQQHGLPLNKADLAVAT
						AKCPISQQQRPILSS
5197	35565	Α	5238	1	483	7.7
5198	35566	Ā	5239	1	1908	
5199	35567	В	5240	1	2718	
	12220,		1:			<u> </u>

SEQ ID NO:	SEQ ID NO: of peptide sequence	4	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5200	35568	Α	5241	522	693	LDFFLVFLQQFLPRPSSSEI*MLP
						GFPAAAYGPVAAAAVAAARGS
						GRKVYGTGDSQA
5201	35569	Α	5242	1	225	MGRNQSRKAENSKHESTYSPP
						KDHSSSQAMEQSWTENGFEKL
						GFRRKEALYYLDLTASCQSQEL
ł						FQKWVSLCCPD*S*TYSPPKDH
						SSSQAMEQSWTENGFEKLGFR
						RKEALYYLDLTASCQSQELFQK
		ļ				WVSLCCPD
5202	35570	Α	5243	1	1129	PWISAPVPVDVVEGAMDSVTV
ŀ						LSFGGLMLYFCAGWPPARRWC
Į						FPESISCGSMERDQWWGLQVA
						KRAGLAGGQSGRTVLRERVRIE
						IASTHIALAARHSDWRCCRNGR
						YPARGPAALQNFQRYTGIQHV
						HRIGMAERMWCDRNRERHTVS
						SSGGNRLPNPGPDRSCDNLKTC
						HTSHGSVMAETAVINHKKRKN
						SPRIVQSNDLTEAAYSLSRDQK
						RMLYLFVDQIRKSDGTLQEHD
						GICEIHVAKYAEIFGLTSAEASK
						DIRQALKSFAGKEVVFYRPEED
						AGDEKGYESFPWFIKRAHSPSR
				-e		GLYSVHINPYLIPFFIGLQNRFT
						QFRLSETKEITDPYAMRLYK\SL
						CQYRAFVNGGGEEKARGKPIL
			<u> </u>			CRYGVGM

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5203	35571	A	5244	1	1753	MVDSLIARVGVMARGNAITLP
		ŀ				VCGRDVKFTLEVLRGDSVEKTS
		ŀ				RVWSGNERDQELLTEDALDDLI
					:	PSFLLTGQQTPAFGRRVSGVIEI
		ŀ				ADGSRRRKAAALTESDYRVLV
						GELHDEQMAALYRLGNDYRPT
		ŀ				SAYERGQRYASRLQNEFAGNIS
						ALADAENISHLHIHAEDTFLPFY
	1					LGKKDDVTYAIKPTCWPGLDII
						PSCLALHGIEIELMGKFDEGKLP
						TDPHLMLGLAIETVAHDYDVIV
				•		IDSAPNLGIGTINVVCAADVLIV
						PTPAELFDYTSALQFFDMLRDL
						LKNVDIKGFEPDVRILLTKYSNS
				Ē		NGSQSPWMEEQIRDAWGSMVL
					•	KNVVRETDEVGKAVINHKKGK
						NSPRMVQSNDLTKAAYSLSRD
	ł					QKRMLYLFVDQIRKSDGTLQE
						HDGICEIHVAKYAEIFGLTSAEA
						SKDIRQALKSLAGKEVVFYRPE
				•		EDAGDEKGYESFPWFIKQSEE/P
						QSGNFYTDTLGMAEFRRGGLR
						ATAGPRLSRTRDCQGTEKVHG
						KTVFDKTVP/WVPKRLKSPSCK
						NRWP*KMQKLSVCTASSPGQQ
						LSTVRVTQREICFPVVRREVLLL
						IGTLTQTRKGQISVRA
5204	35572	A	5245	337	1977	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
5205	35573	I IA	5246	145	1929	 VSGVIEIADGSRRRKAAALTES
3203	333/3		3240	143	1727	DYRVLVGELDDEQMAALSRLG
						NDYRPTSAYERGQRYASRLON
						EFAGNISALADAECDNLKTCHT
	1					SHGSVMAETAVINHKKRKNSP
		ŀ				- · · - · · · · · · · · · · · · · · · ·
						RIVQSNDLTEAAYSLSRDQKRM
						LYLFVDQIRKSDGTLQEHDGIC
						EIHVAKYAEIFGLTSAEASKDIR
		ŀ				QALKSFAGKEVVFYRPEEDAG
		ŀ				DEKGYESFPWFIK/RPSRGLYSV
						HINPYLIPFFIG\LQNRFTQFRLS
						ETKEITNPYAMRLYESLCQYRK
						PDGSGIVSLKIDWIIERYQLPQS
						YQRMPDFRRRFLQGFCRFRNH
						HQTGFSPAGANQRGPLAATLSG
		1				PGGEGQSAVARLTGEKKNHPG
	1	1				AQYANRLSPRVGRFINAAGTTG
			•			FPTWKAGSERNAINDDVTYAIK
						PTCWPGLDIIPSCLALHRIETEL
				·		MGKFDEGKLPTDPHLMLRLAIE
						TVAHDYDVIVIDSAPNLGIGTIN
						VVCAADVLIVPTPAELFDYTSA
						LQFFDMLRDLLKNVDLKGNSN
						GSQSPWMEEQIRDAWGSMVLK
						NVVRETDEVGKGQIRMRTVFE
						QAIDQRSSTGAWRNALSIWEPE
		<u> </u>				CNEISIGVSLDQDGGSNSVLRK
5206	35574	Α	5247	1	1383	
5207	35575	Α	5248	1	1731	
5208	35576	В	5249	1	1290	
5209	35577	В	5250	1	3345	
5210	35578	Α	5251	1	1959	
5211	35579	Α	5252	3	2356	
5212	35580	В	5253	334	1710	

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SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
ļ		1		sequence		
5213	35581	A	5254	1	1302	MKLMETLNQCINAGHEMTKAI
						AIAQFNDDSPEARKITRRWRIG
						EAADLVGVSSQAIRDAEKAGR
						LPHPDMEIRGRVEQRVGYTIEQI
						NHMRDVFGTRLRRAEDVFPPV
1						NVSKSDDTLKINGVEDHKTIFD
						GDGKTYQNVQQFIDEGNYTSG
				İ		DNHTLRDPHYVEDKGHKYLVF
ļ						EANTGTENGYQGEESLFNKAY
						YGGGTNFFRKESQKLQQSAKK
						RDAELANGALGIIELNNDYTLK
						KVMKPLITSNTCDNLKTCHTSH
1						GSVMAETAVINHKKRKNSPRIV
						QSNDLTEAAYSLSRDQKRMLY
						LFVDQIRKSDGTLQEHDGICEIH
						VAKYAEIFGLTSAEASKDIRQA
						LKSFAGKEVVFYRPE/EDAGDE
						KGYESFPWFIKRAHSPSRRLYS
						VHINPYLIPLLYRVPNRVTQFRL
						SETK/EITHPYAMPLYESLCQYS
5214	35582	В	5255	1	1452	
5215	35583	Α	5256	1	1323	
5216	35584	Α	5257	1	1557	
5217	35585	Α	5258	1	2259	
5218	35586	Α	5259	1	2418	
5219	35587	Α	5260	1	1218	MKLMETLNQCINAGHEMTKAI
						AIAQFNDDSPEARKITRRWRIG
						EAADLVGVSSQAIRDAEKAGR
						LPHPDMEIRGRVEQRVGYTIEQI
						NHMRDVFGTRLRRAEDVFPPVI
						GVAAHKERNQNISRYISNRLAR
						RECHDSPEEGTGNTRSKERLQG
						RLPSRGLTESQEHLLDTRKQFW
						FEDFKDGLGPILKCDNLKTCHT
						SHGSVMAETAVINHKKRKNSP
						RIVQSNDLTEAAYSLSRDQKRM
						LYLFVDQIRKSDGTLQEHDGIC
						EIHVAKYAEIFGLTSAEASKDIR
						QALKSFAGKEVVFYRPEEDAG
						DEKGYESFPWFIKRAHSPSRGL
						YSVHINPYLIPFFIGLQNRFTQFR
						LSETKEITNPYAMRLYESLSAS
						NYQRMPDFRRRFLQV\CVMEIN
5220	25500	Ι.	5261	1	2028	KQTSMRLSYI
5220 5221	35588 35589	A B	5261 5262	11	2028 1653	
5222	35590	A	5263	1	1446	
5223	35591	A	5264	1	1788	
		+				
5224	35592	Α	5265	22	1893	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucléotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5225	35593	A	5266		2144	MEKKITGYTTVDISQWHRKEHF EAFQSVAQCTYNQTVQLDITAF LKTVKKNKHKFYPAFIHILARL MNAHPEFRMAMKDGFIENMFF VSANPWVSFTSFDLNVANMDN FFAPVFTMGKYYTQGDKVLMP LAIQFFDMLRDLLKNVDLKGFE PDVRILLTKYSNSNGSQSPWME EQIRDAWGSMVLKNVVRETDE VGKDTGVRSKISGVIKMPMGV AVVNCCTYGSDYRVWLASWM MSQWLHYPEWYRKENKHKFY PAFIHILARLMNAHPEFRMAMK DGELVIWDSVHPCYTVFHEQTE TFSSLWSEYHDDFRQFLHIYSQ DVACYGENLAYFPKGFIENMFF VSANPWVSFTSFDLNVANMDN FFAPVFTMGKYYTQGDKVLMP LAIQFFDMLRDLLKNVDLKGFE PDVRILLTKYSNSNGSQSPWME EQIRDAWGSMVLKNVVRETDE VGKGQIRMGTELEFFRLSETKEI TNPYAMRLYESLCQYRKPDGS GIVSLKIDWIIERYQLPQSYQRM PDFRRFFLQCDNLKTCHTSHGS VMAETAVINHMKRKNSPRIVQ SNDLTEAAYSLSRDQKRMLYL FVDQIRKSDGTLQEHDGICEIHV AKYAEIFGLTSAEASMDIQHAL KSFSGKEVVFYRPEEDAGDEK/ GYESFPWFIKRAHSPSRGLYSV HINPYLIPFFIGLQNRFTQFLLSE
5226	35594	A	5267		1395	MNMNIKKIVKQATVLTFTTALL AGGATQAFAKENNQKAYKETY GVSHITRHDMLQIPKQQQNEKY QVPQFDQSTIKNIESAKGLDVW DSWPLQNADGTVAEYNGYNV VFALAGSPEDADDTSIYMFYQK CDNLKTCHTSHGSVMAETAVI NHKKRKNSPRIVQSNDLTEAAY SLSRDQKRMLYLFVDQIRKSDG TLQEHDGICEIHVAKYAEIFGLT SAEASKDIRQALKSFAGKEVVF YRPEEDAGDEKGYESFPWFIKR AHSPSRGFYSVHINPYLIPFFIGL QNRFTQFRLSETKEITNPYAMR LYESLCQYRK\PDGSGIVSLK/ID WIIKRSQLPQSAFYQPFMGLRR ESFYFRWERRTLGPLKSFSVKR GTEAGKFRLAALLVRL

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5228	35596	A	5269		1375	MNMNIKKIVKQATVLTFTTALL AGGATQAFAKENNQKAYKETY GVSHITRHDMLQIPKQQQNEKY QVPQFDQSTIKNIESAKGLDVW DSWPLQNADGTVAEYNGYHV VFALAGSPKDADDTSIYMFYQ KIRRKNGPVSATFTSDGKIRLFY TDYSGKHYGKQSLTTAQCDNL KTCHTSHGSVMAETAVINHKK RKNSPRIVQSNDLTEAAYSLSR DQKRMLYLFVDQIRKSDGTLQ EHDGICEIHVAKYAEIFGLTSAE ASKDIRQALKSFAGKEVVFYRP EEDAGDEKGYESFPWFIKRAHS PSRGLYSVHINPYLIPFFIGLQNR FTQFRLSETKEITNPYAMRLYES LCQYRKPDGSGIVSL/KIDW/IIE/ RYQLPKVPSPEARKITRRWRI/V KQRI*LGFLLRLSEMPRKQGDY RTRIWKFEDGLSNVLVIQLNKLI ICVMCLVRDCDVLKTYFHR
5229	35597	В	5270	1	1668	
5230	35598	Α	5271	1458	2675	CDNLKTCHTSHGSVMAETAVI NHKKRKNSPRIVQSNDLTEAAY SLSRDQKRMLYLFVDQIRKSDG TLQEHDGICEIHVAKYAEIFGLT SAEASKDIRQALKSFAGKEVVF YRPEEDAGDEKGYESFPWFIKR AHSPSRGLYSVHINPYLIPFFIGL QNRFTQFRLSETKEITNPYAM/R IPLH*LFR*TLRQTKPDNSAGKC VKI**HTQNQRSGRSQNDF*RR GI
5231	35599	В	5272	1	1317	
5232	35600	Α	5273	1	987	
5233	35601	Α	5274	1	571	
5234	35602	A	5275	1	4161	
5235	35603	A A	5276	1	1228	
5236	35604	A	5277	387	3466	
5237	35605	A	5278	1	6721	
5238	35606	A	5279	1	1651	
5239	35607	Α	5280	5082	5879	NULLANDEL TEL COCODOD 4 51
5240	35608	Α	5281	1	231	NHLAVRRLTELSGGQRQRAFL AMVLAQNTPVVLLDEPTTYLDI NHQVDLMRLMGELRTQGKTV VD*VRDLNQVSRYC
5241	35609	Α	5282	1	303	CLRILCPYPYIPLTEKTLA*PAL YTLELTADMAETCKQFLPEFSL TFSFQRGWEKETEYAEVLERNF ERDRQLTYTAHGPHKADLRIQE SGVPYNEILVTL

seq		od	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide
	luence		09/540,217	codon for pentide	L.F tido comunação	
5242 350				• •	or peptide sequence	deletion, \=possible nucleotide insertion)
5242 35				sequence		
	610 A	1	5283	1	273	TNSSVGPGDRLL*QTAAPHRGK
						LKVFFGACAGVGKTWAMLAE
						AQRLRAQGLDIVVGVVETHGR
						KDTAAMLEGLAVLPLKRQAYR
						GRHISEF
5243 35	611 A	1	5284	66	1200	HILPAGHPTLRHSLFPHQQGRSS
		İ				VQSALGAASGNGLKLVGRFEW
	ļ					GPSRRARSGYPDATGYDQ\PKPI
	:			:		SAAVKEFFGSSQLSQFMDQNNP
						LSEITHKRRISALGPGGLTRERA
						GFEVRDVHPTHYGRVCPIETPE
	į					GPNIGLINSLSVYAQTNEYGFLE
						TPYRKVTDGVVTDEIHYLSAIE
						EGNYVIAQANSNLDEEGHFVE
						DLVTCRSKGESSLFSRDQVDY
	ŀ					MDVSTQQVVSVGASLIPFLEHD
	l	į				DANRALMGANMQRQAVPTLR
	ĺ					ADKPLVGTGMERAVAVDSGVS
						GGVVQYVDASRIGIKVTKTRCI
}	ľ					RVKQVSTSTTLPLFLMESNPLF
						WMAVLVSAGILGQWPIGRLAD
						KFGRLLVLRVQVFVVILGSIAM
5244 35	612 E	3	5285	1	862	
5245 35	613 A	4	5286	1	579	DPRD*RLKVAK*MVTLFGPQFV
						RELQQRGFDIFLDLKFHDIPNTA
						AHAVAAAADLGVWMVNVHAS
						GGARMMTAAREALVPFRPDAP
						LLIAVPV*RSIKASDLVDLGMTL
	ĺ					SPADYAERLAALTQKCGLDGV
						VCSAQEAVRFKQVFGQEFKLV
						TPGIRPQGSEAGDQRRIMTPEQ
	Ì					ALSAGVEYMENGRPGTQSVK
5246 35	614 A	4	5287	484	978	GIVRIPHRLADIKCWRAVLYNY
1	j					THYSYQANLCCANLE*ADLSGS
	İ					VLDCANLQGVKMLCSNAEGAS
	i					LKLCNFEDPSGLKANLEGANLK
						GVDMEGSQMTGINLRVATLKN
						AKLKNCNLRGATLAGTDLENC
						DLSGCDLQEANLRGSNVKGAIF
						EEMLTPLHMSQSVR
5247 35	615 A	4	5288	3	504	YRLSQSSSKSLVSRSWLNMRFIS
						SVMFMEPTSSSENRFTSPSCSE
						MSSSEATLRSFFLGTTSPCRSM
						VGLQLVKSSQSSSVSSFSSHVRL
						SQSSSKSKVSRLAGT*EGSEKRS
						FREEKVPCLRLGLPESTLYAHC
I					l l	OV*OPELIA POOTOLII IOOMAAA
	į	- 1				SV*SRTHVFSGTSHHSSKVAVG

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	•			sequence		
5248	35616	A	5289	2	442	HVIWHMPRKTSKCRQLLCSGA
						SRNADTAARQSTCSSHRPPGKIP
						SLGPRRVPGCSSVPSSRGEQSTG
						SPAAPRCGDVTHTEASQEEQP*
						LRGTHGPASTPEQGTAKAKVR
						DRRAVVHPGKTGIPSAAG*KGR
	1					GRPGEPPGAAVQVE
5249	35617	Α	5290	151	1185	YVDQVTHITESPLPSVSYSFAFG
						SLISAVDPVATIAIFNALHVEED
						TEKGHVRLTPASPSTFFFFFSTE
						EKYAPGSCCRKAAVL*LLHISH
						QLSLDSPSLTVVIYVLKHIDLRK
						TPSCRHCEI*VGDHTAVLSVLFS
						PSLGIMAILFSGIVMSHYTPREA
		1				SP*SCCLFTVSSHTVCCHPGLRG
	-	1				AIPYALSLHLDLEPMEKRQLIGT
		1				TTIVIVLFTILLLGGSTMPLIRLM
		}				GLPGGAPRWCQLVAVGPPFLP
		1				AQGNTVESEHLSELTEEEYEAH
		1				YIRRQDQLSLFN*N*SCLWPSPC
						LPPPP*TPHTKHPVASPPFHGPA
						PDALTA WLCLSTQDLHHGRIQ
						MKTLT
5250	35618	Α	5291	5	198	VCGRQQSLEAGL*GAATPFAVF
						PRPWSAVRVKEYGYTRLHILN
						GTHIHIQQVSDDQAFAQCHPP
5251	35619	Α	5292	137	308	HLERTGPNLPPGARLATRPTTR
			ı			LPAASPRGRDT\PLPIPISFYSETE
						AKGEAPYIEPE
5252	35620	Α	5293	612	1078	RPPEKKERGRRVRGRRGGGGK
						RRRKKKKKKGSKGREDAAAA
						AAAAQSPRTRAPAPD*ERRPQP
						APSARAPPRPLLPAPRRSRPAQV
						PPDSASPWAAAAGLALPPPQCP
						GGGARPLRLLPQPPAPGGTAR
						WEPRRPRPAAAPGLLQSPGPLP
		<u>l</u>				RQS
5253	35621	Α	5294	515	817	LALPHLPSNLLLKGWLKLKA*S
						RPSRTPSFG*LSQWKHALKRLK
						PVITCLLQHGFLKPINSPYNSLIL
		1				PVLKPDKPYKLVQDLRLINQIV
						LPIHPMVSQA
5254	35622	Α	5295	1	951	
5255	35623	Α	5296	1	1077	
5256	35624	В	5297	87	1423	
5257	35625	Α	5298	29	234	
5258	35626	В	5299	77	327	
5259	35627	В	5300	1	235	
5260	35628	Α	5301	2	2144	
5261	35629	Α	5302	1	855	
5262	35630	Α	5303	845	1057	

SEQ ID NO:	SEQ ID NO: of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion
5263	35631	A	5304	1	933	
5264	35632	Α	5305	1	954	
5265	35633	A	5306	1	1603	MEAEKFHQMPSVSRRTRKAS
						VIQSKSENLRTRGANSHALKR
					-	KPVITRLLQHGLLKPINSPYNS
						LPVLKPDKPYKRFLAFYGWSI
						SLIVDSCSKLEQHSTLSRVILV
						KGFCRFRNHHQTGFSPAGAN
						RGPLAATLSGPGGEGQSAVA
						TGEKKNHPGAQYANRLSPRV
						RFINAAGTTGFPTGKRAVSAT
						LMDFADFGTTITQDFRLLGQT
	1					VDRLLQLSQGQAVKGNQLLP
						SLVKRKTTLAPNTQTASPRAL
						DSLMQLARQVSRLESGQSSQI
						SLRLTLPDRLRSPLDHHGRRA
	1					GNSHSGSPKPPSCLPLISPHLN
						QVWDISTPSLATDHMPITIPLE
						N\TLTPLNLKCQYPIPQHALKI
						KPVITRVLQHGVLKPTNSPYN
						PILPVLKPDKPYKLVQDLRLII
						IVLLPIHPMVPNPYTLLSSIPPS
						HYSVLDLKHAFFTIPFHPSSQI
						LAFTW\TDPDIHQAQQIT\WA`
						P\QRFPRTGPHYFNQAQISSSS
						TYLGIILIKTYVLSLPIVSD
5266	35634	В	5307	1	874	
5267	35635	A	5308	144	539	DUDA VA GDD GD A GDGGL GA D
5268	35636	Α	5309	1165	1639	PHPAVASRRGRAGPGGLSAPI
						TPSSELSPLSKGDLRKTHRTFS
						Q*GDLRKTHRTFSTR*GDLRK
						HRTFCTR*GDLRKTHRTFSTR
						GDLRKTHRTFSTR*GDLRKTH
		Į			,	TFSTR*GDLRKTHRTFSTR*G
						RKTHRTFCTR*GDLQKTHRTI
						TR*GDLRKTHRTFSTR*GDLR
						THRTFSTR*GDLRKTHRTFCT
		1				GDLRKTHRTFSTR*GDLRKTH
						TFCTR*GDLRKTHRNFSTR*G
						LRKTHRTFSTR*GDLRKTHRT
60.50	105655	4.	5310	03.5	450	CTR*GDLQKTHRTFSTR
5269	35637	Α	5310	235	453	SHWPWLCSSTSCSSQRSIVSSO
						MARGPFLRKRDLLFPPPPPGS
						GLELLEIIFHQGISQVHHLLHL
						VWL*NLNVPDSSLPAENEATO
						GFVPPPLAPVRGPLFPV\D\AR
						FLRKRDLLFPPPPPG\AM\FGA
						DYFPPGDFPGPPPAPFASMAF
	1	1	1			N

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon,/=possible nucleotide deletion, \=possible nucleotide insertion)
5270	35638	A	5311	2	2333	TGRGYCGDHESSFGAMEEPGA TPQPYLGLLEELRRVVAALPE GMRPDSNLYGFPWELVICAAV VGFFAVLFFLWRSFRSVRSRLY VGREKKLALMLSGLIEEKSKLL EKFSLVQKEYEGYEVESSLKDA SFEKEATEAQSLEATCEKLNRS NSELEDEILCLEKELKEEKSKHS EQDELMADISKRIQSLEDESKSL KSQVAEAKMTFQIFQMNEERL KIAIKDALNENSQLQESQKQLL QEAEVWKEQVSELNKQKVTFE DSKVHAEQVLNDKESHIKTLTE RLLKMKDWAAMLGEDITDDD NLELEMNSESENGAYLDNPPK GALKKLIHAAKLNASLKTLEGE RNQIYIQLSEVDKTKEELTEHIK NLQTQQASLQSENTHFENENQ KLQQKLKVMTELYQENEMKL HRKLTVEENYRLEKEEKLSKV DEKISHATEELETYRKRAKDLE
						EELERTIHSYQGQIISHEKKAHD NWLAARNAERNLNDLRKENA HNRQKLTETELKFELLEKDPYA LDVPNTAFGRGSRGPGNPLDH QITNERGESSCDRLTDPHRAPS DTGSLSPPWDQDRRMMFPPPG QSYPDSALPPQRQDRFCSNSGR LSGPAELRSFNMPSLDKMDGS MPSEMESSRNDTKDDLGNLNV PDSSFP\AKKEATGPGFVPPPLA PVRGPLFPVDARGPFLRRGPPFP

SEQ ID NO:	SEQ ID NO: of peptide		SEQ ID NO:	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
5271	35639	A	5312	141	2788	GLRVPLSPSSPRSRLRSGPLRLP
						GADSGSGPKAVCSPPFIVAPTG
						RGYCGDHESSFGAMEEPGATP
						QPYLGLLLEELRRVVAALPEG
						MRPDSNLYGFPWELVICAAVV
						GFFAVLFFLWRSFRSVRSRLYV
						GREKKLALMLSGLIEEKSKLLE
			'			KFSLVQKEYEGYEVESSLKDAS
			ŀ			FEKEATEAQSLEATCEKLNRSN
						SELEDEILCLEKELKEEKSKHSE
						QDELMADISKRIQSLEDESKSL
						KSQVAEAKMTFQIFPMNEERLK
						IAIKDALNENSQLQESQKQLLQ
						EAEVWKEQVSELNKQKVTFED
						SKVHAEQVLNDKESHIKTLTER
						LLKMKDWAAMLGEDITDDDN
						LELEMNSESENGAYLDNPPKG
	ľ					ALKKLIHAAKLNASLKTLEGER
:						NQIYIQLSEVDKTKEELTEHIKN
						LQTEQASLQSENTHFENENQKL
						QQKLKVMTELYQENEMKLHR
						KLTVEENYRLEKEEKLSKVDEK
		1				ISHATEELETYRKRAKDLEEEL
						ERTIHSYQGQIISHEKKAHDNW
		1				LAARNAERNLNDLRKENAHNR
						QKLTETELKFELLEKDPYALDV
						PNTAFGREHSPYGPSPLGWPSS
						ETRAFLSPPTLLEGPLTLSPLLP
						GGGGRGSRGPGNPLDHQITNER
						GESSCORLTDPHRALSOTGFLS
						PPWDQDRRMMFPPPGQSYPDS
	2.5.10	<u> </u>	-	10.15		ALPPQRQDRFCSNSGRLSGPAE
5272	35640	Α	5313	217	447	FQRMSGL*GYSPPLGGQPAL*E
						CQGAADSAGG\RGLDSERSLGR
						\PAASGPNTALGWQAQPWS*Q AGPELPRAEFLQA
5272	25(41	A	5314	3	287	TLRNRHRELRTCLSGPGLGLPT
5273	35641	A	3314	3	207	QVSVVAGGCCVCPSSSQSPVPR
						PPALSAAPWHSHRAGCPPSGGL
		1				*PQRPLILWNQPLSNTLLEKELA
		1				PPPAH
5274	35642	A	5315	3	468	ALARSPAG*PQSPDGDLCFRHR
32/4	33042	^	13313	3	400	KGERRPAAPAGQCPAGRQRHV
						PAAGRDGKPGRAPGADR*AGPI
		1				LHLYHLPAPCGGGTPGDSNPRR
						APAQRSE*CTPPPPQ*RNNRKPS
						TRAGLLLPS*HVQLSGGLTLCP
						GDRVS*WPQGVLAAPGPGPCT
		1				HL
5275	35643	В	5316	64	288	
22/3	33043	ال	12210	194	1200	<u> </u>

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5276	35644	Α	5317	73	265	NGRTSRCFSPGKH*EYCSGTFH
		ŀ				PLQVVGKIYSNLDKIL*ASRAR
						WGLCSRKPGYWSCTQPPTSS
5277	35645	Α	5318	125	461	
5278	35646	Α	5319	298	593	LGVEVEPVTGGDAVIGASHDAF
						LQVNIESIVPEHSTRCKLLGKFI
						RIWIKYFRHLVHGGGSAPVSLV
						TGPVHGDSLLGPVAPEPPRVVE
						PARQVRAHAVVVAGPLRRGRG
				,		LGRHRLRARSG*GMPQSVRGA
						CGAQGHGPAGEDQRLLPRERG
						PAGQVQQPGVVPGLQDPEGCL
		į				RVQDQ*PGLREQSPHRARDA*S
						ILSKFE*IFPTTCSGWNVPEQYS
						QCLPGEKHRDWPR*LHHLL*PA
						QLPPPVTTSTGHPSAIEGLETPL
ŀ						RSPLTPGPPGKSIIPHHPQSPDCD
						GTDCAAFPENIIRLRSRKICKRG
5279	35647	Α	5320	1124	1835	HSLHFPKFPSFLFLTLITLGLLM
						AVPPGLITTHQQRQAML*YKPL
						ARLLEPLISFPSWKSILKEITSQC
						SICYSTTPQRLFRPPPFPTHQTR
						GFAPPR/HWQIDITHMPQVRKL
						KYLLVLVDTGSKKATVVISSLL
						SDIIPRFGLPTSIQSDNGPAFISQI
			ļ			TQAISLALGIQRTHGLLKTHLT
		ļ				NLSHQLKKDWTILLPLSLLRIQ
						ACPRNATGLHSDG/VGPKLLPG
1						SPFQLHHTWDSP
5280	35648	Α	5321	1	1758	
5281	35649	Α	5322	1	1967	
5282	35650	Α	5323	76	1908	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	1	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
5283	35651	A	5324	524	2144	QSDLSASQHGLFPLATEVRSSG
						AASCPDSDSICPAPTIAPGRATP
						PQANCWASEGTLRYQALPGNR
		ŀ			ļ	APVSQVSQAGGANLCVQQPKK
						HLTNFKSGKRPLFTLFSNLQGP
						RSRPVAFLSKQLDLTVLGWPSC
						LRAAAAAALIVLEALKITNYAQ
		ŀ				LTLYSSHNFQNLFSSSHLTHILF
						CSPGSFSCTHSLLSLPQLPIVPGP
						DFSPASHIILDTTPDPHDCISLIH
						LTFTPFPHISFFPAPHPDHA*FIN
						GSSTRPNRHTPAKAGYAIVQVT
						SPPLKTSHFLSIVEIYPQGNNFS
						VFHLLFYYSSGIIQAPLLSLHIKL
ŀ						RDLPPPRTGKSALLNMPRVRKL
		İ				KYLLVLVDTGSKKATVVISSLL
						SDIIPRFGLPTSIQSDNGPAFISQI
		1		,		TQAISLALGIQRTHGLLKTHLT
1						NLSHQLKKDWTILLPLSLLRIQ
		1				ACPRNATGLHSDGEARSSFQAH
						HFNFITHGTVPDVFVPILQLLNE
ŀ		l				DFEDPCLGYLPSPEISICLEPTSW
						SKDRSCSSGEEGSAGQTVTGDL
						HKAQTPELDQPASLNCPDL
5284	35652	Α	5325	1	420	RTRGGEKE*EEKQRSRR*RIK*E
		ŀ				EEEK*RRSRGAGDKG*NERRRR
		ŀ				NRGEPVEKEEDDRRRKRQWRR
		ŀ				GEEEKRRRGRNCGGGGGEEKE
		İ				DGRRRRLPCFLLLPSLLVSLLW
		ŀ				LVSFLPCVFGFFGAPAVVFARF
5005	25.652		5326	2	467	RGLRAFCRF
5285 5286	35653 35654	A	5327	269	777	
5287	35655	A	5328	2443	2627	VGFFFFFWWGERGTGSYSVTQ
3287	33033	^	13320	2443	2027	ARVQ*YDLGSLQLLPPSFKGFS
		1				CLSLPGSWYYRHGPPRLG
5288	35656	A	5329	57	394	obobi do William Mad
5289	35657	A	5330	33	311	HNTLPKTKPWFQEPGDEEPQQE
207						EPPTESRDPAPRLPRSP*SADPP
=						WRSQRPQGLTRPPPEPLRPHGS
						PPAHLSLFCQRSCPGPQERPHPS
	<u>.</u>	1				ТМ
5290	35658	Α	5331	226	406	
5291	35659	В	5332	51	90	
5292	35660	Α	5333	1	348	PPSQRV*GWHEMRQGGRQHAQ
						PVLSQQGLTAGHCHHQHCRRP
						LACDVRPVVYEKVSRDHGLVS
						GQHN/PAPESQGQSTSPPRHTGR
						KQARAGGEKKVTSDPANTAQH
5202	25661	<u> </u> _	5224	100	200	NPNTGSRGRP
5293	35661	Α	5334	109	380	

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/340,217	sequence	of peptide sequence	determiny (possione indetecting insertion)
5294	35662	A	5335	448	724	 WPSRSALAAGGRTAPTPRSRAG
3271	33002	ļ``			,	PPAR*RAGRESARGTS*GIRPRS
		1				QPARRPCPPAPVGRGLPPGPAR
		l				PPPRAAPREPPRCRAAAHTSRP
						AGS
5295	35663	В	5336	1	1554	
5296	35664	В	5337	99	366	
5297	35665	A	5338	1	714	
5298	35666	С	5339	391	689	
5299	35667	A	5340	2	614	
5300	35668	Α	5341	2	461	
5301	35669	В	5342	45	1148	
5302	35670	Α	5343	215	849	IKRLPLMKRMWL*LQPESLEIA
			1			GILVQ*MIE*QPKKGTNSIPVSK
						PSP/VVQKPNGQWRQVQDLRLI
						SDAVIPLYPAVSNPYTLLSQILE
						EAEWFTVLDLKDAFFCIPLRSD
		ŀ				SQFLLAFEDPTDHTSQLTWTVF
						TQGFMDTPHLFGQSLAQDLGH
						FSSPGTLVLQYVDDLLLAKQQA
						TLDLLNFLANQGYKLSKLKAQ
		ļ				LCELLVFSSCARMHS
5303	35671	Α	5344	1	1035	MGQVWALVRSTLELFHTDDEE
						EGEYDEVTEEVTEQVYLPAKA
						KVAQEEEVHPYPSAPPHYYFEE
						KEWPDPPDLSFLEDTGRKVVAP
			1			VTEQHLERLLSVLFRQEFSRLD
			ļ			ERDDAVEQLRGVCIRAWEKITS
						GGEQYPSFSAVKQGPKELYADF
						IAWNLLRQESLKKVISDSAAQD
						IVLQLLAFGNVNLDCQAALRPI
					}	RGKAHLVDYIKACDGIGAKQD
			l			SERFAFTIPVVNNLQPAKHFHY
					•	FTDGSSNGKASYSGSKGQNQQ
						PIWILSRHLKPYHEPDAKEEIPG
						G/CPRTPWLQPCRD*C*GGP*/P
						VTSNTR*TQPPTWGQIKKLSQM
5204	25670	A	5345	1566	1767	VEENLRKAGQLVTMTVYWN KGGWSQRHSQGACYTCRKSGH
5304	35672	A	3343	1300	1707	-
			[WAKECPQPGIPPKPRPICVGPH* KSDCSTHLAANPRAPGTLAQGS
						L
5305	35673	В	5346	1	3429	
5306	35674	Α	5347	1	1776	
5307	35675	Α	5348	1	933	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
5308	35676	A	5349	1	1743	MGKVWAVVRSTLELFHTDDEE
						EGEYNEVTEEITEQVYLPAKAK
İ		ľ				AVKEREQRGNITATFEPFPFKLL
İ		ŀ				KEFKQAINQYGPGSHFVMGLL
	1					KNVAVSSQMIPTDWDALTQAC
	1					LTPAQFLQFKTWWAHEASIQA
	İ					ACNAQAQPQINITTGQLLGVGG
						WAGLDAQVVMQDDAIEQLRG
						VCIRAWEKIPLCGEQYASFSAIK
						QGPKEPYVDFIAWLQESLKKVI
						ADSAAQDIVLQLSAFDNANPDC
1						QAAL*PIRGNAHLVDYIKACDG
:						
	1					IRGNMHKATLLVQAMAGLRVD
						KGNPLLPGACFNCGKHGHTKQ
						ERRKKSVSQAARWGKRKTADP
						EICPKCKKGKYWANQCHSKFD
1						KDGTLISGNAMRAEQDCEWFT
1						FTIPAVNNLQPAKHFHRFTDVS
						SNGKASYSGSKGGQARVQLFE
						NASVRATNSDLPQSSLWCRRKT
						SVSVAVLVSATIPISRVQGPSQV
	ì					LGQGEKQTNQNVVPNPYTLLS
	1					QIPEEAKWFTALDLKDAFFCIPL
						HSDSQFLFAFEDLTDHTSQLTW
						TILPQGFRDSPHLFGQALAQDL
Į.						GHFSSPGTLVLQYVDDLLLATS
						SEASCQQATLDFLSFLAN
5309	35677	Α	5350	1	1085	
5310	35678	Α	5351	<u>l</u>	3105	
5311	35679	Α	5352	1	5172	
5312	35680	Α	5353	41	324	
5313	35681	Α	5354	18	270	KLSQVCIDLL*IREGERDMYPG
						WAKFPSPYSLKGT*PIAQGVLW
						SFGDFFYC/IPSTLPTPKSWRLFE
						SPFPESLTPVSLVWWPC
5314	35682	Α	5355	80	384	SNNRTEGAWGKCQLMSS/PLTE
	1	İ				PQVTLTIDSQEIDFLLDTGAAFS
1						VLISCPEQLSSRSVTI*GILGQPV
}	1					TRYFSHLLSCNWETLLFSHAFL
						VMPESPTPLLGR
5315	35683	С	5356	99	317	
5316	35684	Α	5357	388	476	KVCPRPCRTSM/LPIRII*KLPQV
						CLDLL*IIEGEGDMYPG*AKFPS
						PYCLKGT*PGTIAGSWARGVSA
						AASPSYSSPSY

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
5317	35685	Ā	5358	254	849	LQPESLEIAGILVQ*MIE*QPKK
3317	33003	` `	3330	123.		GTNSIPVSKPSP/GVQKPNGQW
						RQVQDLRLISDAVIPLYPAVSNP
						YTLLSQILEEAEWFTVLDLKDA
						FFCIPLRSDSQFLLAFEDPTDHT
			į.			SQLTWTVFTQGFMDTPHLFGQ
						SLAQDLGHFSSPGTLVLQYVDD
						LLLAKQQATLDLLNFLANQGY
						KLSKLKAQLCELLVFSSCARMH
5318	35686	A	5359	1	909	A SOLD A
5319	35687	A	5360	153	436	CFCCCVEGNKPRNLQRTTKTPR
	33007	` `	3300	1133	1.50	LSVMSPSSCRGRGIWPNPGTC/R
						PSPSLI*SRSRQTWRITTPCQQG
						NKTGQRMNLTN*QEQASEGG*
						*QTPLS
5320	35688	Α	5361	1	1632	4.1.2 5
5321	35689	С	5362	1	675	
5322	35690	A	5363	604	1047	SNNRTDNPTSVAYLSKETDVV
	35070	'				AKGWPHCLWVVVAVAILVLEA
						IKIIQGKDFTVWTSHDVNGILG
						AKGNSWLSDKRLLRYQALLLE
						GPVLQIHTCVALNPAIFLPEDGE
						PIKHDYQQIVAQTYVT*EDLLE
ŀ						VPLANPDLNLYTNGSSFV
5323	35691	Α	5364	749	1002	QLKKGTNSLLVSKPSPVWIPTG
						TLTQIMGTGVANIC*PVF*KD*G
						ELGKMNYAMMSTITQGKEENP
						AFLKWLWEALRKYTPLSP
5324	35692	В	5365	114	2309	
5325	35693	A	5366	413	1157	
5326	35694	Α	5367	105	478	YGHCCYDHRICCFFPATEQKVC
i						PCAAQG*PL*LRGDDPEMSVYI
						*QLPLYYSSDLLSAANEVQILAF
						FQGDVSFFPVATTTDTLSVTFN
						FPFNYQGVNDFDFDFKQFLHSS
					}	FDFCFGRVFSNFE
5327	35695	Α	5368	531	1087	RLQPPGAPVEDITRPVSAKPRD
						ADRNSTGCYLV*LQHQRQTES
						GRQLSGL*GPGCNGVH/ASLYR
						TSAMFAERFGSYSRRSTTAGIPS
						LLRLKSTIRRSDRSRRKAQRKR
1						CGQVMPQHRRSIRPLLQPITPSL
						NRNLHRSPHRSRNRWSRLNRR
		-	}			LKHHLRRSQNRSRSWKKKLHR
İ						RVKPMLCNWVR
	<u> </u>	1	<u> </u>	1	<u> </u>	IN A WE IMPOUM A IC

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	İ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
5328	35696	Α	5369	544	1381	VNPALQVLVVAITPPPSLPSPSCI
						RVLTRVSNAIAAKPLCLAVCAV
						VEGGRIRALVAVAAWGAGVTG
		1				GPAGLIACGPVSPSVIRGQFRHA
		1				GISFFLYAIGGEMVPAVRRONH
						TPA*HKRFAKAFQLAHPNAVN
						AEHFAGGDRHLFSHLNQGCVV
		1				KHHVRRDITFIGYRFTQFA
5329	35697	Α	5370	81	863	AYGNPVEIAGGARQAGVNFTG
	ŀ					VFRPRFSRRWRRISGETCKLWW
				·		RRWRSAWASINQTFASWSTLIF
						RAISNPIIRKPDAPGVMACPRKR
	İ					CCFTIRLIWRGCAVVWKRSR/E
		1				GQLQDIERHKLNAMGAFAEAQ
						TCRRLVLLNYF/GRRASGAVRE
						LRYLPRSAETVRRFNRCSDCPF
						HHWSCESAVWDGLCGGSDSCC
						GVSRARLFASPVNLPITGEIPER
						GFHPFCGRTTENGKNHPRNLFI
						WRSNCSVLPVKVMSLCSSTPCP
5330	35698	В	5371	12	273	
5331	35699	Α	5373	145	336	SEEEEEDEEEEEEEEEEE
	ļ					E*EEGESAIICAKVCISVFPRRK
						NELNMAITPNTIPKLNHW
5332	35700	Α	5374	294	657	VAPGDRHAFHGGGSGLSPELTL
						PQTQCCAQATVQGLEGTRSWS
						QSGTSSLSPWSHTSLACPQKEE
						GEEEEEEEEEEEEEE
	İ					EEEEEEEEEEEEAKTQ*KSG
	<u> </u>					QKI*TDSPQKK
5333	35701	Α	5375	2	588	SRPGITGSTISYGDPGHGAGPPG
						ADLGPARSLKPERAAGAQLAA
		1				GEARQSGDPGLPGGPGHSLGTA
						PC*VDKGWGHPVSTVHHQRQP
						QPGGLRAPGTALLAGTQPSHPA
						AGPWEPQPQRGVRVLGGRRDS
		İ				*VGGG*GQHNKALWGPR*PHT
						EQKPDRKLSRIPLRAAGGEKRG
		<u> </u>				CGCDRAGCLVLGPPQLPAQGL
5334	35702	Α	5376	1	445	SLSHRAGGGGGVGGGAVCLFS
						PPRTKRESAPESPSRVLGHSPVP
						S*GAPSTRGSVSRILTSGGTHER
						SNGLTCHRNQAGTGPSALRSGR
						P*CLSRANPPGSKRSAGPVHEE
						RT*PCRLSGSAGITQ*DGN*KAG
		<u> </u>		<u> </u>	<u> </u>	PRVPEAPERGCATC

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
5335	35703	Α	5377	3	313	QNDEVAFRKLKLIIEDVQGKSY
					İ	LTNFHGMNLTHDKMCSMVKK
						WQTVIEAHVDVKTTNGYLLCL
						FCVGFTKKCNNQIWKTSLF*QQ
						SNMEDFFVLTTGIISHLHEPFLV
						VTDYDQ
5336	35704	Α	5378	45	764	TSFLTTGGKKGS\KKKVVDPFS
						KKD/WL*YVKHRAMFNIRNIGK
						DVGSPRTQGTKIAS\DGLKG\RV
						FE\VSLADLQNDEVAFRKFKLIT
						\EDVQGK\NC\LTNFHGMDLTP*
		1				QKCCSMVKKM/WQTMI*SSPL
						MVKTT\DGLLASVCSCVGFTKK
						\RNNQIRKTSYAQHQQVRQIRK
						KMMEIMTREVQTNDLKEVVNK
						LIPDSIGKDIEKACQSIYPLHDVF
						VRKVKMLKKPKFELGKLMELH
:						GEGS
5337	35705	A	5379	3	501	SSARFVSSSSSGGYGGGYGGVL
	!					TASDGLLAGNEKLTMQNLNDR
						LASYLDKVRALEAAKILSDMRS
		İ				QYEVMAEQNRKDAEAWFTSRT
]						EELNREVAGHTEQLQMSRSEV
						TDLRRTLQGLEIELQSQLSMKA
						ALEDTLAET/EGDVRADSERQN
		1				QEYQRLMDIKSRLEQE
5338	35706	Α	5380	1	716	AQLKVKFWYWPEAGAGGLRR
						FKHY\YTPFKDCRDKILGATIKN
						SRIVLQIDNAGLAADDFRTKFE
						TEQALRMSV*ADITGLRRVLDE
						LTLARTDLEMQIEGLKEELAYL
						KKNHEEEISTLRGQVGGQVSVE
						VDSAPGTDLAKILSDMRSQYEV
						MAEQNRKDAEAWFTSRTEELT
						REVAGHTEQLQMSRSEVTDLR
						RTLQGLEIELQSQLSRLWGFCC
						PLEGVFWVEGWEGRDPYPRLF
5339	35707	Α	5381	3	832	
5340	35708	В	5382	141	1067	

Stop codon, /=possible nucleotide etion, \=possible nucleotide insertion) VAPSVLRLAMTSYSYRQSSAT GFGGLGGGSVRFGPGVAFRAP HGGSGGRGVSVSSARFVSSSS GAYGGGYGGVLTASDGLLAG EKLTMQNLNDRLASYLDKVR LEAANGELEVKIRDWYQKQG GPSRDYSHYYTTIQ\ELRDKIL
VAPSVLRLAMTSYSYRQSSAT FGGLGGGSVRFGPGVAFRAP HGGSGGRGVSVSSARFVSSSS GAYGGGYGGVLTASDGLLAG EKLTMQNLNDRLASYLDKVR LEAANGELEVKIRDWYQKQG
FGGLGGGSVRFGPGVAFRAP HGGSGGRGVSVSSARFVSSSS GAYGGGYGGVLTASDGLLAG EKLTMQNLNDRLASYLDKVR LEAANGELEVKIRDWYQKQG
FGGLGGGSVRFGPGVAFRAP HGGSGGRGVSVSSARFVSSSS GAYGGGYGGVLTASDGLLAG EKLTMQNLNDRLASYLDKVR LEAANGELEVKIRDWYQKQG
HGGSGGRGVSVSSARFVSSSS GAYGGGYGGVLTASDGLLAG EKLTMQNLNDRLASYLDKVR LEAANGELEVKIRDWYQKQG
GAYGGGYGGVLTASDGLLAG EKLTMQNLNDRLASYLDKVR LEAANGELEVKIRDWYQKQG
EKLTMQNLNDRLASYLDKVR LEAANGELEVKIRDWYQKQG
LEAANGELEVKIRDWYQKQG
, ,
ATIENSRIVLQIDNARLAADDF
TKFETEQALRMSVEADINGFR
VLE*S*TLAQEPNLEDARIEG
KEE\LAYLKKNHEEEISTLKGP
CEAQV\SVEVD\SA\PGTRSSPK
SDMRS\QYEVMAEQ\NRKD\A
\\WFT\SR\TERIETREVAWPHR
FQMSQAPRLLDLAAATPQGS
DLSCQSQLEPLKAALGKTPLA
NRKA\RFGNPQLAPYPQALDS
VI*KPQLG/DIVAKLDSSTGRI
EYQ\RL\MDIKFARLEQEIAHP
RSLA/LEGQEDHYNN\LFASK
CLRGSLERCGTLCKI\SLHEISG
ESRVQ\QPLSSREEEEEEEEE
EEEEEEEEEEEEEKK
CSKAQCKHFPLSEVL
EKNEKEQEEEKKEKNSKKK
EEEEEGGGEEEEEGEGGEE
EEEEEEEEEEEEEE
EEE/EEEEEEEEEEEEEE
EEEEEEEEEEEEKKKKK
KKKKKKKKKKKKKKKK
KKKKEEEEKVVEEEDEFIWF
GNSLYFPSFSHFAWSQIPSRED
GERRLRVNSMIMDFTSNTNH
EEEKEEEKEEEKEEEEE
E/DEEGEEGEGEGEEGE
EEEEEEEEEEEEEE
EEEEEEEEEEEEEEE
EEEEEEEEEEEEEEEE
EEEEEEEMIVLKIIVAKQLF
.WLFQEEEEEEEEEED*EEE
EE/EEEEEEEEEEEEEEEE
EEEEEEEEEEEEEE
GHRVGI

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
5351	35719	A	5393		626	EEGEVEEEGEEEEEEGEGEE
3331		``		,	1	EEEEGEGEGEGEGEEEEE
		-				EEEEEEEEEEEEEEEE
	ŀ					EEEEEEEEQEEEEEEEEE
			1			EE\EEAAAAAAASRDSHTSLEL
						TYWPPCPALRDTLVSNCSLSSL
						CTSGDSFHW*ETTIPAMSSCSH
	ŀ					QHSLHQLGIIPRQILMRMASLM
						RRIC\EDHPATAEQ**HV*GPPD
5352	35720	A	5394	2	506	INTERPOLATION OF THE STATE OF T
5353	35721	A	5395	1	480	
5354	35722	A	5396	1	492	
5355	35723	A	5397	1	870	
5356	35724	A	5398	1111	407	
5357	35725	A	5399		651	
5358	35726	В	5400	1	2427	
5359	35727	A	5401	571	966	SPFTNSRLTIFISFSECGQIAEIYI
	33,2,	`				LSQPFMVRIMATEPPIINLQPGN
						FTLDIVSRDHTTAL*PGRQEQNS
						VSKKKEEEEEEEEEEEEE
	1					EEEEEEEEEEEEEEEE
						EEEEEEEEEEEEEEE
						EEEEEEEEEEEEEEKKKK
5360	35728	A	5402	1	1674	
5361	35729	A	5403	1	306	
5362	35730	A	5404	2	431	
5363	35731	A	5405	1	585	
5364	35732	Α	5406	1	708	
5365	35733	A	5407	3	595	
5366	35734	Α	5408	287	659	
5367	35735	A	5409	1	633	
5368	35736	Α	5411	3	634	
5369	35737	Α	5412	344	481	
5370	35738	Α	5413	1	525	
5371	35739	Α	5414	1	501	
5372	35740	Α	5415	1	461	
5373	35741	Α	5416	302	567	MLCCLEISSTRYPKSSLSSSKFH
			}			KSLGQGQNAASLFAKT*RESPL
						LQFPASSSSPSETTSAWTLLFISL
						SAFLTKAFNKSLGGSKLSH
5374	35742	A	5417	251	844	NLGRGSQTSILDFCAPAGLIPCG
						SCQGLGLPHSEATARAVYWPL
						SAMAGVAGTQGTKSLDSSLFA
			1			KT*QESPLLRFPTSSSSPSETTSA
						FILLFISLSEFLSKPFNKSLGGSK
						LSHIFVSSSEPSKLSQPLHMPIQ
					1	ARISSFTSGHTDDSTVLFSVLLK
						LPLPVPSQAFSLKPGSIVMLLSS
						WSDNFNIPSISDSDA
5375	35743	Α	5418	1	1113	
5376	35744	Α	5421	1	337	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5377	35745	Α	5422	2	680	
5378	35746	Α	5423	144	373	
5379	35747	Α	5424	1	597	
5380	35748	A	5425	3	3319	
5381	35749	Α	5426	1	612	
5382	35750	С	5427	46	168	
5383	35751	Α	5428	1	663	
5384	35752 35753	A	5430	67	263	IAIMNDTVAIRTRKFMTNRLLE RKQMVIDVLHPGKATVPKTEIL EKLAQMYKTTPDVIFVFGFRTH CGGGKTTGFGMIYDSLDYAEK NEPKHRLARHGLYEKKKTS*K QRQERTNRMKKVRGTAKANV VAG*KPNE GKTTG\FGMIY\DCLDYAKKNE
						PKHRRARHGLYEKKKTSRKQR KERKNRMKKVRGTAKANVGA GKK
5386	35754	A	5431	2	537	TPGSTTTTKFPTASTSEDRSPIM ND\TVTIPH*EKFMTQPTYFRGN QMVI\DVLSPPGKGDQCPKHRK FGKKLAQNVPRPTPD\VIFVFWI QNFIFGGGKTT\GF\GMIYGFPGI YAKEKMNPDHRLCKDMGLVL RKEKGPSRKANGKGTARNRIG RKVPGGTGKGPNVGCWAKRR NEVSSS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
5387	35755	A	5432	<u> </u>	2195	ARGLSMGH*FP/PHFQCALA*M
3307	33,733	` `		ľ		PAPGRGPPPPLPP/CPDAPPACA
						ATMERPEAGLRVAPRSPVQPGC
						TSVLPVAKRFNARILWLKRPOR
						QLWSLWRAQATVSLGGFHVM
						LSLQVHRINARVKEAWQLPPRF
						QMMYQKAWVLRQKPAAAVEP
	1					SQRNSPRAMLGGNVGLEPPPRV
				•	ļ	STRALPSGAVGRGLPPSRPKND
						TATGSLHPELGKAADIQFHPVR
						AATGASPCKSIGTELPKALGAH
						VLLQCALDVGHGVEGELWSFK
						TRQPSDFSIFSPPFPPFYSTKPPLS
						SWPIPNEPLGTPPRRGRGGAEG
	İ					LLTSQCSHILNGLCFTGGINSNE
						1
	}					CENVSRKKKMSEEFEANTMDS
						LVDMPFATVDIQDDCGIFFDVW
						KLRVNCQEILANRHPGRAMRK
						VVLTKTSWEGESLAVGKLPLK
						RLVSLDCQLRCLQCECGILIRQ
						HVVWLERTTGHWHPRSSQSCR
						LNTTSTALEHCNAFIYNTVCSH
		l				KLKDHSHDNGKSSGFLNPVAA
						PHSCIVMGHLTLLITTSYYTPLT
						KPDTDPTLKKEGLISQDSSSLEV
		1				LLHTDPLGKRGTPDPRVDDDSL
		1				GKFPVTNSQARKRILEPDDFLD
		1				DLDDEYCEDAPKHRGKGCGSA
		1				HKELDASNLEDQDKPYACDIC
		1				GKRYKNRPGLSYHYAHSHLAE
		1				EGEDKEDSQPPTPVSQRSEEQK
		<u> </u>				SKKGPDGFALPNNYCNFCLGNS
5388	35756	Α	5433	1	580	FSPLIHLILSGHSTCFREHRVGG
						KVTDQQDPKAEEFFLVQNKMK
						SLPCLLLSTQTRQPSDFSIFSPPF
						PPFYSTKPPLSSWPVLNELLGTP
						PRRGGGRAEG/PPHFPVG\GPGR
					<u> </u>	GAPSPPRRGGWPGGGLTPQPPS
				ļ		RTGRLAGRGADPPTSLPDGAA
		1				GRAGG*PHHLPPGRGGWPGRG
			<u></u>			APHFPVGVAGQRRLFIC
5389	35757	Α	5434	405	685	KVSHVYFYPHRPRNHPISQFFPH
						PSRLSIPQNRHCHHGPS/RNEPL
						GTPPR/PGSWPGRGAPHFPETFN
						HGRRWKGSRHVFTWPEQERE
						WGSATHF
5390	35758	Α	5435	1	654	
5391	35759	Α	5436	1	426	
5392	35760	Α	5437	1	548	
5393	35761	A	5438	1	459	7.70
5394	35762	Α	5439	1	346	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
5395	35763	Α	5440	1	1248	
5396	35764	A	5441	1	1257	
5397	35765	Α	5442	1	1659	MGDFNTPLSTLDRSTRQKVNK
			:			DTQELNSALHQADLIDIYRSLH
						AKSTEYTFFSAPHHTYSKIDHIL
				1		GSKALLSKCKRTEIITNYLSDHS
						AIKLELRIKNLTQNRSTTWKLN
						NLLLNDYWIHDEMKAEIKMFF
						ETNENKDTTYQNLWDAFKATA
						SKTNKEKEKNQIDTIKNDKGDI
						TTDPTEIQTTIREYYKHLYANK
				}		LENLEEMDKFLDTYTLPRLNQE
						EVESLNRPITGAEIVAIINSLPTK
						KSPGPDGFTAEFYQRYKEELHI
			ĺ			NRAKDKNHMIISIDAEKAFDKI
						QQPFMLKTLNKLGIDGTYFKIIR
						AIYDKPTANIILNGQKLEAFPLK
						TGTRQGCPLSPLLFNIVLEVLAR
						AIRQEKEIKGIQLGKEEVKLSLF
						ADDMTVYLENPIVSAQNLLKLI
						SNFSKVSGYKINVQKSQAFLYT
						NNRQTESQIMSELPFTIASKRIK
						YLGIQLTRDVKDLFKENYKPLL
						KEIKEDTNKWKNIPCSWVGRIN
						IMKMAILPKVIYRFNAIPIKLPM
						TFFTELEKTTLKFIWNQK\RARI
						AKAILSQKNEAGGITLP
5398	35766	Α	5443	1007	1719	TEPKTKTT*LSQ*MQKKPLTKF
						NNPSC*KLSIN/IVLEVLARAIRQ
						EKEIKGIQLGKEEVKLSLFADD
						MIVYLENPIVSAQNLLKLLSNFS
						KVSGYKINVQKSQAFLYTNNR
						QTESQIMSELPFTIASKRIKYLGI
						QLTRDVKHLFKENYKPLLKEIK
						EDTNKWKNIPCSWV\GRINIVK
						MAILPKVIYRFNAVPIKLPMTFF
						TELEKTTLKFIWNQKRACIAKSI
						LSQKNKAGGITLP
5399	35767	В	5444	1	1020	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	1 ' '	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
5400	35768	A	5445	1	1576	MDKFLDTYTLPRLNQEEVESLN
						RPITGAEIVAIINSLPTKKSPGPD
						GFTAEFYQRLISNFSKVSGYRIN
		1				VQES\QAFLYTINRQTESQIMSA
						LPLTIASKRIKYLGIQLTRDVKD
						LFKENYKPLLNEIKEDTNKWK
						NIPCSWVGRINIVKMAILP/KEL
		ļ				EETTLKFIWNQKRARIAKSILSQ
		ļ				KNKAGGITLPDFKLYYKATVT
						KTAWYWYQNRDRDQWNRTEP
				1		SEIMPHIYNYLIFDKPEKNKQW
	1	ŀ				GKDSLFNKWCWENWLAICRKL
						KLDPFLTPYTRINSRWIKDLHV
						RRKTIKTLEENLGNTIQDIGMG
						KDFMSKTPKAMATKAKIDKW
						DLIKLKSFCTAKETTIRVNRQPT
						KWEKIFATYSSDKGLISRIYNEL
						KQIYKKKTNNPIKKWAKDMNR
		1				HFSKEDIYAAKKHMKKCSSSLA
		l				IREMQIKTTMRYHLTPVRMAII
		1				KKSGNN/R/CAPGTPERQNHSLS
						WKGS*SQEPSGLAQWIPLPWSP
						AS*DPLA*NSCCQHSCLKSTWD
						AQACARKLRTLIKGYRNCQLE
5401	35769	Α	5446	1	984	
5402	35770	Α	5447	1	1563	
5403	35771	A	5448	1	1449	
5404	35772	В	5449		1605	
5405	35773	В	5450	1	1569	MCCI DCTIACKBURYI CIQI TDD
5406	35774	Α	5451	1	690	MSELPFTIASKRIKYLGIQLTRD
						VKDLFKENYNPLLNEIKEDTNE WKNIPCSWVGRINIVKMAILPK
	}					1
						ATVTKIAWYWYQNRDIDQWN RTERSEIMPHIYNYLIFKKPEKN
						ſ
						KQWGKDSLFDKWCSENWLAIC RKLKLDPFLTPYTKINSRWIKD
						LNVRPKTIKTLEENLGITIQDIGI
						GKDFMSKTPKAMATKAKIDK
						WDLIKLK\FCTAKETTIRVNRQS TKWEIISSLD
5407	25775	 	5452	1	1919	I K WEHOOLD
5407	35775	Α	5452	1	1717	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
·	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
·				sequence		
5408	35776	Α	5453	11	1416	MIILIDAEKAFDKIQQPFMLKTL
						SKLGTDGTYLKIIRAIYDKPTAN
						IILNGQKLEAFPLKTGTRQGCPL
		ĺ				SPLLFNIGLEDLARAIRQEKEIK
						GIQLGKEEVKLSLFADDMIVYL
						ENPIVSAQNLLKLISNFSKVSGY
			:			KINVQKSQAFLYTNNRQTESQI
		l	:			MSELPFTIASKRIKYLGIQLTRD
		l				VKDLFKENYKPLLKEIKEDTNK
		1				WKNIPCSWVGRINIVKMAILP\R
		l				DIDQW/NRTEPSEIMPHTYNYLI
		l				FDKPEKNKQWGKDSLFHKWC
						WENWLAVCRKLKLDPFLTPYT
						KINSRWIKDLNIRPKTIKTLEEN
						LGITIQDIGVGKDFMSKAPKAM
						ATKAKIDKWDLIKLKSFCTAKE
						i i
į						TTIRVNRQPTTWEKIFATYSSD KGLISGIYNELKQIYKKKTNNPI
Ì		1				KKWAKDMNRHFSKEDIHAAK
			}			KHMKKCSSSLAIREMDIKTTMR
		l				YHLTPVRMAIIKKSGNNRCWR
		1				GCGEIGTL
5409	35777	В	5454	1	810	GCGEIGTE
5410	35778	В	5455	1	1277	
5411	35779	В	5456	130	1701	-
5412	35780	A	5457	1	1395	
5413	35781	Α	5458	3	1646	
5414	35782	Α	5459	1	759	
5415	35783	A	5460	2821	5781	
5416	35784	В	5461	1	2025	
5417	35785	Α	5462	1	3663	
5418	35786	В	5463	1	2265	
5419	35787	Α	5464	1	2241	
5420	35788	Α	5465	1	2478	
5421	35789	Α	5466	1	2067	
5422	35790	A	5467	1	2988	
5423	35791	В	5468	1	2061	
5424	35792	Α	5469	1	2141	
5425	35793	Α	5470	1	400	
5426	35794	A	5471	2	2507	
5427	35795	Α	5472	1	768	
5428	35796	Α	5473	1	1527	
5429	35797	Α	5474	1	1293 •	
5430	35798	A	5475	1	1038	
5431	35799	A	5476	1	1572	
5432	35800	Α	5477	11	1290	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
5433	35801	Α	5478	11	524	MSELPFRIASKRIKYLGIQLTRD
						VKDLFKENYKPLLNEIKEDTNK
	1	ļ				WKNISCSWVGRINIVKMAILPK/
						VKHR/FSACLQSLGRSGTTGLE
						ALAM*PIWPQEVE*VKFPALPS
						GASSTKGGCVP*QIQVEERLLS
			ļ			WKL*QA*TSW\YQRQRRSDSSL
						RERK*PKSIQRNVRRLSYYTP
5434	35802	Α	5479	2	1547	
5435	35803	Α	5480	1	2376	
5436	35804	Α	5481	1	1824	
5437	35805	Α	5482	1	1071	
5438	35806	В	5483	1	1839	
5439	35807	В	5484	1	1368	
5440	35808	Α	5485	1	1884	
5441	35809	Α	5486	1	1053	
5442	35810	В	5487	1	2532	
5443	35811	Α	5488	1	2373	
5444	35812	Α	5489	1	2361	
5445	35813	Α	5490	1	2274	
5446	35814	Α	5491	1	1686	MASPGPAPTASPKIHSELLTATR
	ŀ					EAQRHHPVPRGQDLVTSESFSL
					ļ	SFCFSAAIFIFELLGSNSEGVTDL
						RLWLCQPAPRCGEWTYNPLEQ
	1					CCDDGVILDLNQTRLCGSSCTF
						WPCFQHCCLESLGSQNQTVVRF
						KVPGMKPDCKSSPITRICAQAG
						VQISNFSKVSGYKINVQKSQAF
						LYTNNRQTESQIMSELPFTIASQ
						RIKYLGIQLTRDVKDLFKENYK
						PLLKEIKEDTNKWKNIPCSWVG
						RINIVKMAILPKVIYRFNAIPIKL
						PMTFFTELEKTTLKFIWNQKRA
						LIAKSILSQKNKAGGITLPDFKL
						YYKATVTKTAWYWYQNRDID
						QWNRTEPSEITPHIYNYLIFDKP
						EKNKQWGKDSLFNKWCWENW
						LAICRKLKLDPFLTPYTKINSRW
						IKDLNVRPKTIKTLEENLGITIQ
					1	DIGMSKDFMSKTPKAMATKDK
						IDKWDLIKLKSFCTAKETTIRV\
						NRQPTKWEKIFATYSSDKGLIS
	1					RIYNELKQIYKKKTNNPIKKWA
						KDMNRHFSKEDIYAAKKHMK
1						KCSLSLAIREMQIKTTNQNH\M
-						RYHLTPGHG
5447	35815	Α	5492	1	1872	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5448	35816	A	5493	195	1494	PPGQHLQRRCLLRVSGFSENAK KGMLEVLARAI\RQEKKIKGIQS GKEEVKLSLFADDMIVYLENPI VSAQNLLKLISNFSKALGYKISV QKSQAFLYTNNRQTESQIMSEL PFTIASKRIKYLGIQLTRDVKDL FKENYKPLLNEIKEDTNKWKNI PCSWIGRINIVKMAILPKVIYGF NAIPIKLPMTFFTELEKTTLKFI WNQKRARIAKSILSQKDKAGGI TLPDFKLYYKATVTKTACNSN GSQSPWMEEQIRDAWGSMVLK NVVRETDEVGKGQIRMRTVFE QAIDQRSSTDTSLSTPAAPMVD SLIARVGVMARGNAITLPVCGR DVKFTLEVLRGDSVEKTSRVW SGNERDQELLTEDALDDLIPSFL LTGQQTPAFGRRVSGVIEIADG SRRRKAAALTESDYRVLGGEL DDEQMAALSRLGNDY
5449	35817	Α	5494	1	1845	
5450	35818	Α	5495	1	2109	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5451	35819	A	5496	l	2535	MKSGHPEKEQDNSDVQETREIT
						IRGLLCTALMRHSTGAIAYLGV
						LSGSASLKLAGVPLRCCEGDKD
						AGHPLETQTALCERGRGARSLV
						GNTIMTSQPVPNETIIVLPSNVIN
						FSQAEKPEPTNQGQDSLKKHLH
						AEIKVIGVNLIQNVLERGWGKC
						QEMIYVLGLDICRPFFVSRVSEE
						GRMGQRGEEDANSLDFPPASLL
						CLICQEQGVNGESCSPVGMYH
						REIVPVYEVLSVITGLQIQVFSG
						KEADSVIKRSIGWGPFFKPRTK
						DKNHMIISIDAEKAFDKIQQHF
						MLKTLSKLGIDGTYLKIIRAIYD
						KPTANIILNGQKLEAFPLKTGTR
	1					QGCPLSPLLFNIVLEVLARAIRQ
	,					EKEIKGIQLGKEEVKLSLFADD
		:				MIVYLENPIVSDQNLLKLISNFS
						KVSGYKINVQKSQAFLYTNNR QTESQIMSELPFTIASKRIKYLGI
1						QLTRDVKDLFKENYKPLLNEIK EDTNKWKNIPCSWVGRINIVK
						MAILPKVIYRFNAIPIKLPMTFF
						TELEKTTLKFIWYQKRARITKSI
		İ				LSQRNKAGDITLPDFKLYYKAT
						VNKTAWYWHQNRHIDQWNRT
						KPSEITLHIYNYLFFDNPDKNKK
		1				WGKDSLFNKWCWENWLAICR
						KLKLDPFLTPYTKINSRWIKDL
						NIRPKTIKTLEENLGITIQDIGMG
						KYFMTKTPKAMATKAKIDKW
		ŀ				DLIKLKSFCTGKETTIRVNRQPT
5452	35820	A	5497	3	2284	
5453	35821	A	5498	1	1245	
5454	35822	A	5499	1	1140	
5455	35823	A	5500	383	1185	
5456	35824	Α	5501	1	3159	MLEVLAWAVRQEKEIKGIQLG
						KEEVKLSLFADNMTVYLENPIV
		ŀ				SAQNLLKLISNFSKVSGYKVSG
		İ				QKSQALLYTNNRQTESQIMSEL
		ļ				PFTIASKRIKYLGIHLTRDVKDL
		ŀ				FKENYKPLLKEIKKDTNKWKNI
						PCSWVGRINIVKMAILPKDIIQE
						NFPNLARQANIQIQEIRKTPQRY
						SSRRATPRHIIVRFTKVEMKEK
						MLRAAREKASHHTYSKIDPILG
						SKPLLSKCKRTEIITNYLSDHSAI
						KLEFRIKNL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
ĺ	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
5457	35825	Α	5502	583	2515	MQKKPLTKFNNPSC*KLSIN*LI
						SNFSEVSGYKISVQKSQAFLYT
	}					NNR/QESQIMSELPFTIASKRIKY
						LGIQLTRDVKDLFKENYKPLLK
						EIKEDTNKWKNIPCSWVGRINI
į		ł				VKMAILPKAIYRFNAIPIKLPMT
	1				ĺ	FFTELEKTTLKFIWNQKRARIA
		l				KSILSQKNKAGGITLPDFKLYY
		l				KATVTKTAWYWYQNRDIDQW
						NRTEPSEITPHIYNYLIFDKPEK
,	1					NKQWGKDSLFNKWCWENWLA
						ICRKLKLDPFLTPYTKINSRWIK
						DLNVGPNTIKTLEENLGITIQDI
						GTGKDFMSKTPKAMATKAKID
						KRDLIKLNSFCTAKETTIRVNR
						QPTKWEKIFTTYSSDKGLISRIY
						NELKQIYKKKTNNPIKKWAKD
						MNRHFSKEHIYAAKKHMKKCS
						PSLAIREMQIKTTMRYHLTPVR
						MAIIKKSENNSLLAAGGGNRRT
						ANVVAHGFANLLTLDKKTLQEI
		İ				LVHYPDSERILMKKARVLLKQ
						KAKTAEATPPRKDLALLFPPKE
						ETPKLFKTLLGGTGKASLARLL
						KLKREQAAQKKENSEGGEEEG
						KENEDKQKENEDKG
						KENEDKDKGREPEEKPLDRPEC
						TASPIAVEEEPHSVRRTVLPRGT
		1				SRQSLIISMAPSAEGGEEVLTIE
5458	35826	Α	5503	1	2463	
5459	35827	В	5504	1	1221	
5460	35828	Α	5505	1	4494	
5461	35829	Α	5506	1	2825	MARTSYGQHFFPTLISMNIDSG
						RQNNGIPKLSTSYSLKPMNTLC
		ŀ				YLVKRDLEGVLKLRILTWESSL
						DYPGGPSVITKVLIRQKEGQSQ
						RTRYDNGSRGWSDAIAVKRPQ
		1		1		AKECGWPLEAGKGKEWILRKE
						CIPANTLMLAQCPPTLRVGGSV
						CSAWRMIQNSDGERRKRNRIQL
			-			QAVRLSKVIHDTFRNFPRFVHS
						RELSKGAEKSSSLKKGLEQVII
						WTKCRQCGESRCTKKVEERKV
		1		1		LHNTFISTSTTITTLG
L			L	J	<u> </u>	LIMITIOTOTITITU

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
5462	35830	A	5508	1184	4791	TSMQKSSIKYWQTESSSTSKSLS
						TMIKWASSLGCKAGSVYANQ*
						M*SSI*TEPKTKTI*LSQ*MQKK
						PL/YKIQQPFMLKTLNKLAQNL
		1				LKLISNFSKVSGYKINVQKSQA
						FLYTNNRQTESQIMSELPFTIAS
		1				KRIKYLGIQLTRDVKDLFKENY
						KPLLKEIKEDTNKWKNIPCSWV
						GRINIVKMAILPQVIYRFNAIPIK
						LPMTFFTELEKTTFKFIWNQKR
						ARIAKSILSQKNKAGGITLPDFK
5160	25021	ļ.,	5500		2022	LYYKATVT
5463 5464	35831 35832	A B	5509 5510	516 286	2233 4131	
5465	35833	В	5511	1	4911	
5466	35834	A	5512	1080	3119	SSGLHPWDARLVQYTQINKCN
15400	155054		3312	1000	3117	PAYKQSQRQKPHDYLNRCRKK
ļ		ŀ				AFDKIQQPFMLQTLNKLGIDGT
						YFKIIRAIYDKPTANIILNGQKLE
						AFPLKTGTRQGCPLSLLLFNIVL
						EVLARAIRQEKEIKGIQLGKEEV
						KLSLFADDMIVYLENPIISAQNL
						LKLISNFSKVSGYKINVQKSQTF
						LYNNNRQTESQIMSELPFTIALK
						RIKYLGIQLTRDVKDLFKENYK
						PLLKEIKEDTNKWKNIPCSWVG
						RINIVKMAILPKVIYRFNAIPIKL
						PMTFFTELEKTTLKFIWNQKRA
						RIAKSILSQKNKAGGITLPDFKL
		ŀ				YYKATVTKTAWYWYQNRDID
		ŀ				QWNRTEPSEIMPHIYNYLIFDKP
						EKNKQWGKDSLFNKWCWENW
		İ				LAICRKLKLDPFLTRYTKINSR
						WIKDLNVRPKTIKTLEENLGITI
						QDIGVGKDFMSKTPKAMAAKA
		Ė				KIDKWDLIKLKSFCTAKETTIRV
					1	NRQPTTWEKIFATYSSDKGLISR
		1				IYNELQQIYKKKTNNPIKKWAK
						DMNRHFSKEDIYAAKKHMKK\
						CSSSLAIR\EMQIKTTMRYHLTP
						VRMAIIKKSGNNRIRRHFQPRS AMFFCCCPGGEGSGLQWELLL
						-
		1				AAWTVLPYPCPAATVGRFLKQ
						AQTADHSFLGHVWRVPSSRSW
						TDKEILSLVVNRQNQEVVQLIG
		<u></u>	l	1		GLCGTHTISPTQHPATG

SEQ ID	SEQ ID NO: Met SEQ ID NO: Nucleotide				Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
			į	sequence		
5467	35835	Α	5513	2494	4549	LQTILQGYSNSKQHGTGTKTA
						WYWYQNRDIDQWNRTEPSEIT
						PHIYNYLIFDKPDKNKKWGKDS
						LFNKWCWENWLVICIKLKLDP
					•	FLTPYTKINSRWIKDLNVRPKTI
						KTIEENLGNTIQDIGMGKDFMS
						ETLKARATKAKIDKWDLIKLKS
						FCTAKETTIRVNRQPTKWEKIF
						AIYSSDKGLIPRIYKELKQIYKK
						KTNSPIKKWVKDMNRHFSKEDI
						YAAKKHMKKCSSSLAIREMQI
						KTAMRYHLTPVRMAI
5468	35836	Α	5514	1	5582	MGDFNTPLSTLDRSTRQKVNK
						DTQELNSALHQADLIDIYQTLH
				1		PKSTEYTFFSAPHHTYSKIDHIL
						GSKALLSKCKRTQIITNYLSDHS
						AIKLELRIKNLTQNHSTTWKLN
						NLLLNDYWVYNEMKAEIKMFF
						ETNENEDTTYQNLWDAFKAVC
						RGKFIALNAHKRKQERSKIDTL
						TSQLKELEKQEQTHSKASRRQE
						ITKIRAELKEIETQKTLQKINESR
}						SWFFERINKIDRPLARLIKKKRE
		l				KNQIDTIKNDK
5469	35837	Α	5515	3371	5375	TDTSQKKTFMQPKKHMKKCSP
						SLAIREMQIKTTMRYHLTPVRM
						AIIKKSGNNRSWYFEKINKIDRL
						LARLIKKKREKNQIDAIKNDKG
						DITTDPTEIQTTTREYYKHLYA
						NKLENLEEMDKFLDTYNLPRL
						KQEEVESLNRPIRGSEIVAIINSL
						PTKKSPGPDGFTAEFYHRYKEE
				ľ		LVPLLLKLFQSIEKEVILPNSFYE
						ASIILIPKPDRDTAKKENFRPISL
						MNIDAKILNKILANRIQQHIKKL
		<u> </u>				IHHDQVG

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5470	35838	A	5516	,	2695	GSACRDPDPDPLRLVGGGAAA PPCSSVSPSAWRQSRPREYYEG GRGLHAPDRETALGLGIATSER MLGICRGRRKFLAASLSLLCIPA ITWIYLFSGSFEDGKPVSLSPLE SQAHSPRYTASSQRERESLEVR MREVEEENRALRRQLSLAQGR APSHRRGNHSKTYSMEEGTGD SENLRAGIVAGNSSECGQQPVV EKC\RNPLHFHLIADSIAEQILA TLFQTWMVPAVRVDFYNADEL KSEVSWIPNKHYSGIYGLMKLV LTKTLPANLERVIVLDTDITFAT DIAELWAVFHKFKGQQVLGLV ENQSDWYLGNLWKNHRPWPA LGRGYNTGVILLLLDKLRKMK WEQMWRLTAERELMGMLSTS LADQDIFNAVIKQNPFLVYQLP CFWNVQLSDHTRSEQCYRDVS DLKVIHWNSPKKLRVKNKHVE FFRNLYLTFLEYDGNLLRRELF GCPSEADVNSENLQKQLSELDE DDLCYEFRRERFTVHRTHLYFL HYEYEPAADSTDVTLVAQLSM DRLQMLEAICKHWEGPISLALY LSDAEAQQFLRYAQGSEVLMS
						RHNVGYHIVYKEGQFYPVNLL RNVAMKHISTPYMFLSDIDFLP MYGLYEYLRKSVIQLDLANTK KAMIVPAFETLRYRLSFPKSKA ELLSMLDMGTLFTFRYHVWTK GHAPTNFAKWRTATTPYRVEW

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
5471	35839	Ī _A	5517	13	2079	GKPVSLSPLESQAHSPRYTASS
	33037	ľ	331,			QRERESLEVRMREVEEENRALR
						RQLSLAQGRAPSHRRGNHSKT
						YSMEEGTGDSENLRAGIVAGNS
	İ	ŀ				SECGQQPVVEKC\RRNPLHFHLI
						ADSIAEQILATLFQTWMVPAVR
		1				VDFYNADELKSEVSWIPNKHYS
						GIYGLMKLVLTKTLPANLERVI
						VLDTDITFATDIAELWAVFHKF
						KGQQVLGLVENQSDWYLGNL
						WKNHRPWPALGRGYNTGVILL
						LLDKLRKMKWEQMWRLTAER
						ELMGMLSTSLADQDIFNAVIKQ
						NPFLVYQLPCFWNVQLSDHTRS
	1					EQCYRDVSDLKVIHWNSPKKL
		ŀ				RVKNKHVEFFRNLYLTFLEYD
						GNLLRRELFGCPSEADVNSENL
•						QKQLSELDEDDLCYEFRRERFT
				:		VHRTHLYFLHYEYEPAADSTD
•						VTLVAQLSMDRLQMLEAICKH
						WEGPISLALYLSDAEAQQFLRY
			}			AQGSEVLMSRHNVGYHIVYKE
	1		ŀ			GQFYPVNLLRNVAMKHISTPY
						MFLSDIDFLPMYGLYEYLRKSV
						IQLDLANTKKAMIVPAFETLRY
						RLSFPKSKAELLSMLDMGTLFT
	}					FRYHVWTKGHAPTNFAKWRT
	1					ATTPYRVEWEADFEPYVVVRR
ŀ						DCPEYDRRFVGFGWNKVAHIM
						ELDVQEYEFIVLPNAYMIHMPH
						APSFDITKFRSNKQYRICLKTLK
						EEFQQDMSRRYGFAALKYLTA
5472	35840	Α	5518	403	2757	
5473	35841	В	5519	128	353	
5474	35842	Α	5520	2	333	
5475	35843	В	5521	25	1359	
5476	35844	В	5522	1	1113	
5477	35845	Α	5523	1	2526	
5478	35846	Α	5524	3	555	

SEQ ID	ISEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
5479	35847	IA	5525	1	1710	I MSQGTTPWGPTPAGTTPESEVH
34/9	33047	A	3323		1710	RPFPFPVPVPVPETSERSFPVRG
						VCGQAKSGLRRGAARLAPAAY
						LFTRQQTRELIHSACVSISVNTM
		l				HGLSVAINSRRCTYYRKYVIDF
						SKSEIIGIPISKNIDLTVGTTVTQ
		1				LQNLNTVGIIGSRGGRGQVAAL
						NCDRQGHLWLFTRVTVHWRM
		1				GNDRTFWGLLDTVSELMFIPGN
				1		PKHHCGPPVKVGAYGVTFLGV
						HWCGACQDTPCKMKDKLLHL
						APPTTKKKAQYVVGLLGFWRPI
						YRVTQKAASFEWGPEQDKALP
						QVQAAVQPALLFGPYDSADPM
						VLEASVADRDAVWSLWQAP/IS
						HKMGHAQQHSIIKWK*YICDW
						ARAGPKGTT\APMASWGVLYD
						QLTEEEKTRAWFTDGSARYAG
						TTQKWTAAALQPLSRTSLKGS
						GEGKSSQWAELQAVHLVVHFS
						WKDKWPDVRLYIDSWAVANG
						LAGWSGTWKKHDWKIGDKEI
				1		WGRGMWMDLSEWPKPVKIFG
]		SHVSAHQWVISAEEDFNNQVD
						KMTCSVDITQPLSPATPVITQW
						AHKQSGHGGRDGGYTWAQQH
						GLPLTKTGLAMATAECPI
5480	35848	Α	5526	1	438	-
5481	35849	Α	5527	l	612	
5482	35850	Α	5528	2	736	EKCAMTALSSKLISQQKAFFAK
ļ						MVVDAVMMLDDLLQLKMIGI
		l				KKVQGGALEDSQLVAGVAFKK
1						TFSYAGFEMQPKKYHNPKIALL
						NVELELKAEKDNAEIRVHTVED
						YQAIVDAEWNILYD\KLEKIHH
						SGAKVVLSKLPIGDVATQYF\A
						DRDMFCAGRVPEEDLKRTMM
						ACGGSIQTSVNALSADVLGRCQ
	1					VFEETQIGGERYNFFTGCPKAK
İ						TCTFILRGGAEQFMEETERSLH
						DAIMIVRRA
5483	35851	Α	5529	3	474	
5484	35852	Α	5530	374	968	
5485	35853	Α	5531	1	568	
5486	35854	Α	5532	1	2136	
5487	35855	Α	5533	1	240	
5488	35856	Α	5534	248	1034	
5489	35857	Α	5535	1	1602	
5490	35858	Α	5536	120	340	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5491 35859	35859	A	5537	2	408	DEDDDDDDEEDDDEDDDDDF DDEEAEEKAPVKKSIRDTPA\K NAQKSNQNGRDSKP\SSTPRSK
						GQESFKKQEKTPKTPK\GPSSVE
						DIKAKMQASIEKGGSLPKVEAK
						FINYVKDCF\RMTDQEAI\QDLW QWRKSL
5492	35860	Α	5538	194	1027	WSFISIHLLSYLSACHHLMEDS/
						MDMGMRPLRPQNYLFSCELKA
						NKDYHFKVDNNENEHQLPLTTI
						NLGAGAMDELDIVEAEAMNYE
						GSPIKVILATLKMSVQPTVSLG
		ŀ			ŀ	G\FEITPPVV\LRLKCGSGPVHIS
:						GQHLVAVEEDAESEDEEEEDV
		ľ				KLLSISGKRSAPGGGSKVPQKK
		ŀ				VKLAADEDDDDDDEEDDDED
		ŀ				DDDDDFDDEEAEEKAPVKKSIR
		ŀ				DTPAKNAQKSNQNGKDSKPSS
		ļ				TPRSKGQESFKKQQKTPKTPKG
						PSSVEDIKAKMQASIEKAH
5493	35861	С	5539	212	367	
5494	35862	Α	5540	14	1515	
5495	35863	Α	5541	1	540	
5496	35864	Α	5542	1	1881	MDAALDDLIDTLGGPEETEEN
						TTYTGPEVSDPMSSTYIEELGK
 -		İ				REVTIPPKYRELLAKPIGPDDAI
		ŀ				DALSSDFTCGSP*LYPSQVFHLL
		ŀ				DLLPFSPGYQPRLLQHHLA/AVP
					İ	FKVAGSGGADCILHMEVRDTA
•		İ				SDTGAGAAALDSSVFSVGLDGF
		ŀ				SLHSDRSKSSESSSMSSDSLRGL GFSSGSGNSGFPSLSVAGFNLYS
		ŀ				DGIVSSSSPHFSSFSSLAFASSTS
		ŀ				LIERRSSSGSACRVPSEADRASS
5497	35865	Α	5543	1	930	LIERRSSSUSACKVFSEADRASS
5498	35866	A	5544	1	183	
5499	35867	A	5545	1	555	
5500	35868	A	5546	162	499	FPGSGNMAKDAGLIEANGELK
						VFIDQNLSPGKGVVSLVAVHPS
						TVNPLGKQLLPKTFGQSNVNIA
		ŀ				QQVVIGTPQRPAASNTL/VWVG
		ŀ				SPHTPSTHFASQ\NQPSDSSPW\S
}						AGKR
5501	35869	A	5547	2	451	LVAEFSAADNHILPNESAYDQK
		ł	1			NIRRRVYDALNVLMAMNIISKE
						KKEIKWIGLPTNSAQECQNLEV
						ERQRRLERIKQKQSQLQELILQ
		.				QIAFKNLVQRNR\HAEQQASRP
						PPPNSVIHLPFIIVNTSKKTVIDC
						SISNDKFEYLFNFDNT
5502	35870	Α	5548	37	162	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
5503	35871	IA	5549	205	616	RKLHGSQSSITIAVFGKQNTYIR
5505	33671	^	3347	203		LEPFKINVLEQITKHIEKLQCGG
		1				VVKQLSRRGNNQHISSTYDINR
		ŀ				
						ADAQVRRAVNNYDIIVMSNSF
						NGQSEHQVWIGGQFTFIKF\PIN
		L		<u> </u>	1.405	SVSIRCRARQEGIMSSPGQQVG
5504	35872	В	5550	<u> </u>	1407	
5505	35873	Α	5551	3	416	MLPARMCCGIVSG*QFHTGNSY
						DHDYEFELGTRS*A*LKLLFS*T
						LPD*QLPAGSGEGPYHLEGQLS
		•	:			YCHRGGEKALAALLSSPTSKTR
	<u> </u>					SPSEPDPEQDEQKLRFCKRHLY
						GQQPRSPVEIRLQHVAIAYQTH
		l				HAYD
5506	35874	Α	5552	1988	2137	LVITTLLPLACGTAK*E*VNVTS
						FGSRPICSTSPVLFSGL*GPVKEF
						рт
5507	35875	В	5553	1	561	
5508	35876	Α	5554	1	492	
5509	35877	Α	5555	3	1055	
5510	35878	A	5556	1	634	
5511	35879	Α	5557		723	
5512	35880	A	5558	li	384	
5513	35881	A	5559	1	792	
5514	35882	A	5560	3	1607	HCTRMSVKWTSVILLIQLSFCFS
	33002	ľ.				SGNCGKVLVWAAEYSHWMNI
						KTILDELIQRGHEVTVLASSASI
						LFDPNNSSALKIEIYPTSLTKTEL
						ENFIMQQIKRWSDLPKDTFWLY
<u> </u>						FSQVQEIMSIFGDITRKFCKDVV
						SNKKFMKKVQESRFDVIFADAI
						FPCSELLAELFNIPFVYSLSFSPG
						YTFEKHSGGFIFPPSYVPVVMSE
						LTDQMTFMERVKNMIYVLYFD
						FWFEIFDMKKWDQFYSEVLGR
						PTTLSETMGKADVWLIRNSWN
						FQFPHPLLPNVDFVGGLHCKPA
						KPLPKEMEDFVQSSGENGVVV
						FSLGSMVSNMTEERANVIASAL
						AQIPQKVLW\RFDGD\KPDTLGL
						NTRLYKW\IPQN\DLLGHPKTRA
						FITHGGANGIYEAIYHGIPMVGI
		1				PLFADQPDNIAHMKARGAAVR
		1				VDFNTMSSTDLLNALKRVINDP
		1				SYKENVMKLSRIQHDQPVKPL
						DRAVFWIEFVMRHKGAKHLRV
		1				AAHDLTWFQYHSLDVIGFLLV
[1				CVATVIFIVTKCCLFCFWKFAR
		1				KAKKGKND
5515	25002	ļ	55(1		1600	NANNUNINU
5515	35883	Α	5561	2	1600	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
5516	35884	A	5562	2	1624	AATGKQALHCIRMSMKWTSAL
3310	33664		3302	2	1021	LLIQLSCYFSSGSCGKVLVWPT
						EFSHWMNIKTILDELVQRGHEV
		Ī				TVLASSASISFDPNSPSTLKFEV
						YPVSLTKTEFEDIIKQLVKRWA
				:		ELPKDTFWSYFSQVQEIMWTFN
	Ì					DILRKFCKDIVSNKKLMKKLQE
	į					SRFDVVLADAVFPFGELLAELL
	1					KIPFVYRPRFSPGYAIEKHSGGL
						LFPPSYVPVVMSELSDQMTFIE
				;		RVKNMIYVLYFEFWFQIFDMK
	1					KWDQFYSEVLGRPTTLSETMA
	1					KADIWLIRNYWDFQFPHPLLPN
						VEFVGGLHCKPAKPLPKEME\E
		1				FVQSSGENGVVVFSLGSMVSNT
						SEERANVIASALAKIPQKVLWR
						FDGNKPDTLGLNTRLYKWIPQ
						NDLLGHPKTRAFITHGGANGIY
						EAIYHGIPMVGVPLLADQPDNI
						AHMKAKGAAVSLDFHTMSSTD
					:	LLNALKTVINDPLYKENAMKLS
						RIHHDQPVKPLDRAVFWIEFVM
						RHKGAKHLRVAAHDLTWFQY
:						HSLDVTGFLLACVATVIFIITKC
						LFCVWKFVRTGKKGKRD
5517	35885	A	5563	1	225	
5518	35886	Α	5564	1	3210	
5519	35887	Α	5565	1	835	HDD CLODICH KVI TKKVI KKVI
5520	35888	A	5566	222	477	HPRCLSPKGILKYLTKKYLKKN
						NLRDWLRVVANSKESYELRYF
						QINQDEEEEEDED*ISFIWKILYE
5501	2.5000	ļ. —	5567	1	770	FLNKTWDPKKKKTKKKKK MAAAAAAAGDSDSWDADAFS
5521	35889	Α	5567	1	778	
						VEDPVRKVGGGGTAGGDRWG
						GEDEDEDVKDNWDDDDDEKK
		l				EEAEVKPEVKISEKKKIAEKIKE
						KERQQKKRQEEIKKRLEEPEEP
:		1				KVLTPEEQLADKLRLKKLQEES
						DLELAKETFGVNNAVYGIDAM
						NPSSRDDFTEFGKLLKDI\ITQYE
						ESLYYARFLEVLVRDVCISLEID
						DLKKITNSLTVLCSEKQKQEKQ
						SKAKKKKKGVVPGGGLKATM
		<u> </u>				KDDLADYGGYDGGYVQDYED
5522	35890	A	5568	1	897	
5523	35891	A	5569	1	2658	
5524	35892	A	5570	1	441	
5525	35893	A	5571	3	427	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
5526	35894	Α	5572	1	952	PRLVLLKMAPLDLDKYVEIARL
						CKYLPENDLKRLCDYVCDLLL
						EESNVQPVSTPVTVCGDIHGQF
						YDLCELFRTGGQVPDTNYIFMG
						DFVDRGYYSLETFTYLLALKAK
						WPDRITLLRGNHESRQITQVYG
		ļ				F\YDDCLTKYG\NANAWRYCTK
			ļ			VFDMLTVAALIDEQILCVHGGL
		l				SPDIKTL\D\QIRTIERNQEIPHKG
		İ				AF\CDL\VWS\DPEDVDT\WAI\S
						PRGAGWLFGAK\VTN\EFGS*ST
		l				NLKLICRAHQLVHEGYKFMFD
						EKLVTVWSAPNYCYRCGNIASI
						MVFKDVNTREPKLFRAVPDSE
						RVIPPRTTTPYFL
5527	35895	A	5573	54	588	
5528	35896	Α	5574	1	4287	
5529	35897	Λ	5575	1	507	
5530	35898	Α	5576	1	1416	
5531	35899	A	5577	1	841	ILLWDVDGGLTQIDKYLYSSED
						YIKSGALLACGIVNSGVRNECD
		ŀ				PALALLSDYVLHNSNTMRLGSI
						FGLGLAYAGSNREDVLTLLLPV
		İ				MGDSKSSMEVAGVTALACGMI
						AVGSCNGDVTSTILQTIMEKSE
						TELKDTYARWLPLGLGLNHLG
						KGEAIEAILAALEVVSEPFRSFA
						NTLVDVCAYAGSGNVLKVQQL
						LHICSEHFD\SKEKEEDKDKKEK
						KDKDKKEAPADMGAHQGVAV
						LGIALIAMGEEIGAEMALRTFG
						HLLRYGEPTLRRAVPLALAL
5532	35900	A	5578	2	391	AFGVTEPGCYGVIDVDTGKSTL
						FVPRLPASHATWMGKL/HSVLT
						SQKPSVLLTLRGVNTDSGSVCR
						EASFDGISKFEVNNTILHPEIVE
		1				CRVFKTDMELEVLRYTNKISSE
						AHREVMKAVKVGMKEYELER

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5533	35901	A	5579	18	1541	RRCRANMAAATGPSFWLGNET LKVPLALFALNRQRLCERLRKN PAVQAGSIVVLQGGEETQRYCT DTGVLFRQESFFHWAFGVTEPG CYGVIDVDTGKSTLFVPRLPAS HATWMGKIHSKEHFKEKYAVD DVQYVDEIASVLTSQKPSVLLT LRGVNTDSGSVCREASFDGISK FEVNNTILHPEIVECRVFKT\DM EL\EVLRYTNKISS\EAHREV\MK AVKSGEWKEYGVGKASFEHY\ CYSRG\GMRH\TSYT\CILRAVG *GTSAVLTFTGH\AGAPNDRTIQ NGDMCLFDMGGEYYSVASDIT CSFPRNGKFTADQ\KAVYEAVL LSSRAVMGAMKPGDWWPDM\ HRLAYRIHLEELAHMGILSGSV DAMVQAHLGAVFMPHGLGHF LGIDVHDVGGYPEGVERIDEPG LRSLRTARHLQPGMVLTVEPGI YFIDHLLDEALADPARASFLNR EVLQRFRGFGGVRIEEDVVVID
						SGIELLTCVPRTVEEIEACMAGC
		L.				DKAFTPFSGPK
5534	35902	A	5580	1	98	
5535	35903	Α	5581	1	810	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	:	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
•				sequence		
5536	35904	Α	5582	1	861	MLKEIIDDTNKWKHIPCSWMG
						RISIVKMTILPKAIYKFNAVLIKI
						PPSFFTELEKTILKFLCNEKRAS
						RAKARLNKKNKSGGITLLDFKR
						YYSAIVTKTVWYWYKNRHIHO
		l				RNRIENPEINPNTYSQLIFNKAN
						KNIKWGNDTLFSKWCWDNWL
						ATCRRMKLDPHLSPHTKINSRW
						IKDLILRPETTKILEDNIGKTLLD
	1					VGLGKDFMTKNPKANARKAKI
						NIWDLIKLKSFFTAKGTVSRVN
						RQPTEWEKIFIIHTSDEGLLISRI
						1
						YNELKQISKKK\TNNPIKKWAK
						DMNRQFSKEDVQMANKYMKK
						CSTSLIANLICREMQIKTTMQY
						HLTPARMAIIKKSKISRCWHGC
						GDQGTLLHCWWECKLVQPLW
		[KTVWRFLKELKVELPFDPAIPL
						LGTHPEEKKSLCKKDTCTCMLL
		1				AAQFTIAKSWNQPKCPSVNEWI
	1	ĺ				KKLCPRWLLSKWASA*NKWK
						HIPCSWMGRISIVKMTILPKAIY
						KFNAVLIKIPPSFFTELEKTILKF
		l				LCNEKRASRAKARLNKKNKSG
		l				GITLLDFKRYYSAIVTKTVWY
	}	1				WYKNRHIHQRNRIENPEINPNT
	1	1				YSQLIFNKANKNIKWGNDTLFS
	1	ļ				KWCWDNWLATCRRMKLDPHL
		1				SPHTKINSRWIKDLILRPETTKIL
	1	1				EDNIGKTLLDVGLGKDFMTKN
		1				PKANARKAKINIWDLIKLKSFF
		1				TAKGTVSRVNRQPTEWEKIFIIH
5537	35905	В	5583	473	1694	
5538	35906	Α	5584	1	429	
5539	35907	Α	5585	572	1149	PPVSHSMTLGNSYLFAPNWTIT
						PAW/CNTLDPSEGFGWQASLSL
						PSHQQFEHTCPEWLPTYSWPCK
						VGPDYLSEICQPECKQHFQESL
						TESSKQNIYETCAIYVSSPGERIL
	1					NQRFGPHGAFILVREYDDIRFS
						HSGPPSTCPNFPGPSYYTARESA
						YALKLRALEIIQISQSTKMQRNL
				-		GNPILPAIHQQPPQ
5540	35908	A	5586	1	804	
5541	35909	В	5587	1	1995	
••	1 <u>-</u>		1,	L	1 - ·	<u> </u>

SEQ ID	SEQ ID NO:		SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
5542	35910	A	5588	74	1128	LLLDL*VD*SPSRPLEMRFRV*P
3342	33710	1	3300	, -	1120	MKRCTTLKKNYLSFLVYISRNL
						*NRVKAIM/VGKAEWKTIELPL
						PRKTVNQKQYRIPGGIAEISTTI
						KELNNAGMVIPTTSPFNSPIWP
						VQKIDGSWRMTVDYCKLNQV
						VTPIAATVPYVVSSLEQIN/TSPG
}						T/WNRKRFCNRSRLLHKLLCHL
						GHMTQQI*CFLRCQWHIGMLF
						GPEICGCTSLSGQKL/WKIFVSH
						AASFEWSPEQEKALQQVQAVV
						QAALPLGPYDPADPVLLEMSV
						ADRDAVWILWQAPISESQWRPP
						GFWSKALPSSADNYSPFERQLL
						ACYWTLVETERLTMGHQVTIPP
						ELPIMNWVFSDLSSPKMCHPQQ
5543	35911	A	5589	1	1337	MIISIDAEKAFDKIQQPFMLKTL
3343	33911	A	3369	1	1557	SKLEETASPSPVVATYTPQPML
						PSAFPPLSEEINPVLPETTVMAS
						PEAVTRQDNVDSPQKPPPTPMF
						ASRPITRLKPRRAPSEEGIQRLK
ŀ						KIGMVEWISHFRPTPLSMEGPE
						HILLTNTLLNRYVKAAPASLKS
-						PLTALLFMSDLTVGTTFSQLQN
						LNTMGIFGSSCDRSQVAALNHQ
						RQVPECKIVIDILNIWRIPHIGSL
						TGRVREIMVGKAKWKPLEQPP
						PRKIVIQKQYHIPGAITEISATIE
						DLKETGVVILTTSPFKSSIWPLQ
						KTDGSWRMTVNYHKLNQVET
						PIAAAVPDVVSLLEQINTSPGT
]		WYAAIDLVNVFFSIPVYKAHQK
						QFAFSWQGQQYTFTFLPQGYIN
						SPDITLVHYIDDIMLIGSSEQEV
						ANTLDLLPARVASWGVPYDQL
		ļ	,			T/GGRED*GLLHRWSCRICRHH
5544	35912	_	5590	2	693	
3344	133912	Α	المودد	4	093	KIVN/QKQYHIRGGIAEISATIKD SKDTGVVIPTTSPFNSPIWPVW
						KTDGSWTITVDYHKLDQVVTPI
1						,
1						AAAVPDVVSLLEQINTSPGTWY
						AAINLENAFFSIPVHKANQ/KQF
						AFSWQGQQYTIAVLPQ\VKFLG
						IHWFVACQDIPSKIKDKLLHLA
1						PPTNKKEAQCLVGLFGFWRQHI
-						PHLRVLLQPIYQVTLKAAIFEW
						GPEQEKALQQVQAAVQAALPL
L		l				GPYDPTDPMVLE

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5545	35913	I A	5591	11	1011	MVSTPATLPSLPKPALMASWG
		ĺ				VPYDQLTKEEKTRVWFTDGSA
						RYAGTTQKWTAVALQPLSRTS
	ļ					LKDSSEGKSSQWAELQAVYLV
						VHFAWKEKWPDVGLYTDSWA
					ł	VANGLAGWSETWEKQDWKIG
:						DKEIWGRGMWMDLSEWSKAV
						KIFVSHVSAHQRVTSAEEEFNN
						QVDRPL/PVFTQWAHEQSGHSG
						RDGGYSWAQQT\GLPFTKADL
						AMATAECPICQQQRPTLSPLYS
						TIPQGDQPATWWQIDYIGPLPS
						WKGQKFVLTVIDTYSRYRFAY
	ł					PAHNASAKTTIHGLIECLIHCYG
						IPHSIASDQSIHF/TTKEVQ*WAH
						AHGIHWSYHVSHHPEAAGL
5546	35914	A	5592	7	278	SASCTAAFHHQMEVVYTRSGS
3346	33914	A	3392	1'	278	SRS*R\PTIPVIAQWDYDHNGHG
						GRDGGYPWAQQHELPLTKADL
	1					ATATAECPICQQQRPSLSPRCGT
						IPR
5547	35915	A	5593	1	237	IF K
5548	35916	C	5594	63	116	
5549	35917	A	5595	2	739	HKMGHAQQHSIIKWK*YICDW
	33717	'	3373	[,3,	ARAGPKGTT\APMASWGVLYD
	1	ı				QLTEEKTRAWFTDGSARYAG
						TTQKWTAAALQPLSRTSLKGS
						GEGKSSQWAELQAVHLVVHFS
						WKDKWPDVRLYIDSWAVANG
						LAGWSGTWKKHDWKIGDKEI
						WGRGMWMDLSEWPKPVKIFG
						SHVSAHQWVISAEEDFNNQVD
		1		:		KMTCSVDITQPLSPATPVITQW
						AHKQSGHGGRDGGYTWAQQH
ł						GLPLTKTGLAMATAECPI
5550	35918	A	5596	1	354	
5551	35919	A	5597	3	592	
5552	35920	A	5598	126	365	QLAEPPHWLPD*KGAGFEWGP
						EQKKALQ*VQAAVQAALPLGP
						YDPADPMVLVVSVADKDAVWI
						FHLSGSDRWRRTYRCL
5553	35921	A	5599	1	483	
5554	35922	Α	5600	1	1908	
5555	35923	В	5601	141	1569	
5556	35924	В	5602	46	1533	
5557	35925	Α	5603	61	448	WRFLGPWRIVSCWFCGWEERR
						EGSAFFVGIPARMSAARESHPH
						GVKRSASPDDDLGSSNWEAAY
						LGNEERKQKFLRLMGA/GKGEE
						DQKINEELESQYQHSMDSKLSG
l	1				1	RYRRHCGLGFSEVEDHDGEGD

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
5558	35926	I A	5604	<u> </u>	404	 PWRIVSCWFCGWEERREGSAFF
3336	33920	^	3004	'	404	VGIPARMSAARESHPHGVKRSA
						SPDDDLGSSNWEAADLGNEER
						1
		1		ļ		KQKFLRLMGAGKKEHTGRLVI
						GDHKSTSHFRTG\EDHDGEGDV
						AGDDDDDDDDDSPDP*SPDDSES DSESE
5559	35927	Α	5605	2	434	LGPWRIVSCWFCGWEERREGS
3333	33727	l'`	3003			AFFVGIPARMSAARESHPHGVK
						RSASPDDDFFLFAIMNGK*KIV
						NSHRCSVKAKSRQQLGSSNWE
		ł				AADLGNEERKQKFLRLMGAGK
		1				KEHTGRLVIGDHKSTSHFRTGE
		1				ED*KINEELESQYQH
5560	35928	С	5606	909	1096	ED KINEELESQ I QH
5561	35928	A	5607	1	309	PQPAAMAAYIWVLIRHGESAW
3301	33929	A .	3007	1	1509	NLETSFSGWYDATLSLAGHKE
						AKRGGQALRGLFQVPLAPEAV
		l				GTVWCSWRTPCALKQL*PRPL
						KRRLRILADRHATQPRPQQ
5562	35930	Α	5608	93	914	SQSVPHPQPAAMAAYK\LVLG\
3302	33930	^	0000	33		RHGESAWNLENRFSGWDDADL
						SPAGQRGGERFAAQALQDAGY
						EFDICFTSV\QKRAIRTLW\TVL\
		ŀ				DAFDQMWLASGEGLWGFNER
						HYWGS*PGLN*KQKLAAKIG*G
						PRLKI\WRRSY\D\VPPPPMEPDH
]
						PFYSNISKDRRYADLTEDQLPS
						C\ESLKDTIARALPFWNEEIVPQI
						KEGKRVLIAAHGNSLRGIVKHL
						EGLSEEAIMELNLPTGIPIVYEL
						DKNLKPIKPMQFLGDEETVRKA
5.50	0.5001	<u> </u>	5.600		450	MEAVAAQGKAKK
5563	35931	Α	5609	2	450	VNKAGGLIYQLDSYAP/RAEAE
						KTFSYPLDLLLKLHDERVLVAF
						GQRDGIRVGHAVLAINGMDVN
						GRYTADGKEVLEYLGNPANYP
						VSIRFGRPRLTSNEKLMLASMF
						HS\IKFVVLADP\RQAGIDSLLRK
		 				IYEIYSDFALKNPFYSLEMP
5564	35932	A	5610	1	663	
5565	35933	Α	5611	1	2034	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
5566	35934	A	5612	374	1106	IPVSERQGSDWRFFSVYVVNKA
			,			GGLIYQLDSYAPTGLRLEKTF\S
						YPLDLAGSS*HDERVLVAFGQR
		1				GRHPEWGHAVL\AINGM\DVNG
						RYTADG\KEVLEYLG*PLLNYP
			ļ			VSIRF\GRPRL\TSNE\KLMLGLP
						CFHSLFAIG\SQACLPGTREASR
						HLKILGRQDTFK\LH\CYQTLT\G
	ŀ					IKFVVL\ADPR\QAGI\DSLLRKI
						YE\IY\SDFALKNP\F\YSLEMPIR
						CELFDQNLKLALEVAEKAGTF
5567	35935	Α	5613	1	336	
5568	35936	В	5614	60	2175	
5569	35937	Α	5615	2	198	
5570	35938	Α	5616	1	449	
5571	35939	Α	5617	1	771	
5572	35940	Α	5618	18	736	EAMSSKVSRDTLYEAVREVLH
						GNQRKRRKFLETVELQISLKNY
						DPQKDKRFSGT/VQVGTVLIPPS
					ŀ	PQCPRACPSPAGSR*AGGGHVG
			:			TDVLG*FRPPAPGRRGWTDPHP
						GS*NMRGGVGRPRPSPPASLRS
						QAGCW*M*TLRVRLGWLLRSQ
						TLHRALGTPGLGVTCWCCPPK
						RRLKSTPRPKFSVCVLGDQQHC
						DEAKAVDIPHMDIEALKKLNK
						NKKLVKKLAKKYDAFLASESLI
5573	35941	Α	5619	24	698	EAMSSKVSRDTLYDAVREVLH
						GNQRKRRKFLETVELQISLKNY
						DPQKDKRFSGTVRLKSTPRPKF
						SVCVLGDQQHCDEAKAVDIPH
			:			MDIEALKKLNKNKKTGSRSWP
						KKYDAFLA/SQSPLIQADSPKSL
						GP\SLNKGRKSSPSLLTH\NKNM
						VAKVDEVKSTIKFPNWRRCLCL
						AVAVGHVKMTDDELVYNIHLA
						VNFLVSLLKKNWQNVRALYIK
						SPMGKPQRLY
5574	35942	Α	5620	1	843	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5575	35943	Α	5621	3	1215	LLIMADPRDKALQDYRKKLLE
		l				HKEIDGRLKELREQLKELTKQY
						EKSENDLKALQSV\GQI\VGEVL
		1				KQLT\EEKFIVKATNGPRYVVG
	1					C\RRQL*QKVSLKPGTR\VAL\D
	1				ĺ	MTTLTI\MEDILPR\EVDPLVYN
	1	1				MS\HEDPG\NVSYSEIGGLSE\QI
						RGI*GEVIGLPLYKPQSYFQRVG
						IIPPK\GCLLYG\PPGTGKNT/LLA
	}					RAVASQLDCNFLKVVSSSIVDK
						YIGESARLIREMFNYARDHQPCI
						IFMDEIDAIGGRRFSEGTSADRE
						IQRTLMELLNQMDGFDTLHRV
						KMTMATNRPDTLDPALLRPGR
		ŀ				LDRKIHIDLPNEQARLDIL\KIHA
						GPITKHG\EIDYEAIVKLSDG\FN
		Ì				GADLRNVCT\EAGMFAIRADHD
						FVSTGKTSLKAV\RKVADSKKL
		l				ESKLDYKPV
5576	35944	Α	5622	104	173	
5577	35945	Α	5623]	1140	
5578	35946	Α	5624	1	528	
5579	35947	A	5625	1	1083	
5580	35948	Α	5626	1	471	
5581	35949	Α	5627	1	2322	
5582	35950	Α	5628	1	678	
5583	35951	В	5629	213	1477	
5584	35952	Α	5630	2	339	FGLHKWGLGLEASSPCGPLTPS
						YLGFPRPHPGTLGLGRFWPFLA
						LQSLSETPSHARMPRVTVARTS
-						PETLEISPLRAAT*NHRSSGHTC
						FRERMLRVSCRLSLDLSSSWVR
5585	35953	Α	5631	1	783	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
5586	35954	A	5632	92	2746	ASRMTVLQEPVQAAIWQALNH
						YAYRDAVFLAERLYAEVHSEE
						ALFLLATCYYRSGKAYKAYRL
						LKGHSCTTPQCKYLLAKCCVD
	1					LSKLAEGEQILSGGVFNKQKSH
	ļ					DDIVTEFGDSACFTLSLLGHGH
						VQTERAAKGTAFSPKRLCLHTF
						P\WTPLEPLVEIGERPLPHQTFN\
						FTS*RNFANCLHTSGTKQVTN\
			:			HVFSHRQPETVLTETPQ\DTIEL
						NRLNLESSNSKYSLNTDSSVSYI
1	1					DSAVISP\DTVPLGTG\TSILSKP
						VQ\NKPKTGRSLLGGPAALSPL
						TPKFWGFLPIRKPP\SPG\DGS\Y
						LQNYTNTPPVIDVPSTGAPSKK
						TFRVLQSVARIGQTGTKSVFSQ
						SGNSREVTPILAQTQSSGPQTST
						TPQVLSPTITSPPNALPRRSSRLF
				1		TSDSSTTKENSKKLKMKFPPKIP
						NRKTKSKTNKGGITQPNINDSL
						EITKLDSSIISEGKISTITPQIQAF
						NLQKAAAEGLMSLLREMGKG
						YLALCSYNCKEAINILSHLPSHH
						YNTGWVLCQIGRAYFELSEYM
						QAERIFSEVRRIENYRVEGMEIY
						STTLWHLQKDVALSVLSKDLT
						DMDKNSPESGTQRGRESPRMC
						GADSQIEKSQSSYFIQDLSRFDY
						VKFAVVTKIRAWCAAGNCFSL
						QREHDIAIKFFQRAIQVDPNYA
						YAYTLLGHEFVLTEELDKALAC
						FRNAIRVNPRHYNAWYGLGMI
5587	35955	A	5633	260	678	GEFSIFFWVWN*LLLWHQETFS
						I*NLFSSH*TSITYQATEEGFDFII
						QMKLIHFGDSIFIYLKLLILNRK
1						P**EAAISI*V*TWRSRFSHFQLR
						*HQTQVTNHSWFTEAR*LTDLV
						KSKSPTIPCSRFKANTLF**SDP
5588	35956	A	5634	1	1452	
5589	35957	Α	5635	2	1563	
5590	35958	A	5636	1	444	
5591	35959	Α	5637	2	345	

SEQ ID NO:	SEQ ID NO: of peptide	Met	SEQ ID NO:	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
110.	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
5592	35960	A	5638	2	1721	RSREAAAVAAAAATTAFGCRI WNPCAALTMKQSSNVPAFLSK LWTLVEETHTNEFITWSQNGQS FLVLDEQRFAKEILPKYFKHNN MASFVRQLNMYGFRKVVHIDS GIVKQERDGPVEFQHPYFKQGQ DDLLENIKRKVSSSKPEENKIRQ EDLTKIISSAQKVQIKQETIESRL SELKSENESLWKEVSELRAKHA QQQQVIRKIVQFIVTLVQNNQL VSLKRKRPLLLNTNGAQKKNL FQHIVKEPTDNHHHKVPHSRTE
						GLKPRERISDDIIIYDVTDDNAD EENIPVIPETNEDVISDPSNCSQY PDIVIVEDDNEDEYAPVIQSGEQ NEPARESLSSGSDGSSPLMSSA VQLNGSSSLTS\EDPV\TMDGFP FGMDNIQSFWGRVEL\LDY\LDS IDCSLEDFQAMLSGRQFSIDPDL LVDLFTSSVQMNPTDYINNTKS ENKGLETTK\NNVVQPVSEEGK ENLKSKP\DK\QLIQYTAFPLLAF LDG\NPASSV*TGRGTTASFRKF CPL*DKPIEVDELLDSSLDPEPT QSKLVRLEPLTEAEASEATLFY LCELAPAPLDSDMPLLDS
5593	35961	Α	5639	2	251	
5594	35962	A	5640	70	297	QLRSPLKWKKAQRQPLAIAHV LLLRPRKKRQNGFFTVRYARLL STLPRSWRR\KQWY*AQNHVRS PEWKWHYQSLS
5595	35963	Α	5641	3	985	
5596	35964	Α	5642	1	3501	
5597	35965	Α	5643	1	1752	
5598	35966	С	5644	1	879	
5599	35967	A	5645	21	519	LHIPVRTLSCRFVGVCWRSTPG PVCLGISSGGCRTVDIAALQLW LRGLALPEQLLRVDPHPQACRR LPREPP\GPGQTFGTREPPQAVS TKEASSSNLHAPERTVAGLTFT TEQVRALEGVFRHHQYLGPLE RNWLAREMQLS\RSR*KPGFKI AG*NTNGKCRTPS
5600	35968	Α	5646	90	2401	
5601	35969	A	5647	776	1018	
5602	35970	В	5648	1	1296	
5603	35971	Α	5649	l	1257	
5604	35972	Α	5650	731	4841	
5605	35973	Α	5651	325	9497	
5606	35974	Α	5652	1	322	
5607	35975	Α	5653	1	819	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5608	35976	Α	5654	56	251	
5609	35977	A	5655	149	701	
5610	35978	Α	5656	3	415	
5611	35979	A	5657	3	467	AIASPRAAGIRHELTSTMAAGK NKRLTKGGKKGAKKKAV/DNII NIGKTLVTRTQRTKIASDGLKG RVFEESLADLQND\TDGYLLRVI *VAFTTERTNQI/REVFNKLIPDS IGKDIEKACQSIYPLHDDFARK
5612	35980	A	5658	2 .	418	VKMLKKPKFELRKLMELHGEG PRVRADGKNFRLTKGGELGAR NKVVDPFSKKDWYDVKAPAM FNIRNIGKTLVSRTQGTKIASDG LKGRVFEARLADLQNDEVAFIK CKLITDDVHGKNCLTIFHGMDL TRDKM\CSLGKKWQTMIEAHV DVKTTDGYLL
5613	35981	Α	5659	3	121	
5614	35982	Α	5660	1	237	
5615	35983	A	5661	25	967	FSPAAGIRHEGSLTSTMAVGKN KRLTKG\GKKGS\KKKVVDPFP R\KDWY*RRKHPLMFHYKN\IG KT\LVTQDPKGPKIA\SDGLQGF VCLKVSLADLQ\NDEVASRKF\ RLITEVCFRGKNLP*LTSHGAW DLYPVTKMCSHGQKNWQT\MI DSSR*MFKTNPMGYLALRLFLC /VGFT*KNRNQFRLRKAPPYAS ATQQVPPKSRKKM\ME\IMTPR GARQNDLEKKWVNKLIPRQAL GKDIRKGPAQSYFIPL\HDVLRL EKLKLLEGSPKF*IGESSLGASM GGRAVVPGKSHLGTQTGAKVG TSLMGYEPPVPRILV

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5616	35984	Α	5662	1	2142	MIILIDAEKAFDKIQQPFMLKTL
			İ			NKLGIDGTYLKITRAIYDKPTA
						NIILNGQKLEAFPLKTGTRQGCP
						LSPLLFNIVLEVLAQAIRQEKEI
						KGIQLGKEEVKLSLFADDMILY
						LENPIVSAQKLLKLISNVSKVSG
		1				YKINVQKSQAFLYTNNRQTESQ
		1				IMSEFPFTIATKRIKYLGIQLTRD
		ŀ	1			VKDLFKKYKPLLNKIKEDTNK
		1				WKNIACSWIGRINIMKMAFPR
						WELNNENTWTQEGEHHTLGPV
		İ				VGWGKRGGIALVDIPNVNDKL
						MVLEVLARAIRQKKEIKGIQLG
						KEEVKLSLFADDMIVYLENSIV
						SAQNLKLISNFSKVSGYKINVQ
		ļ				KSQAFLYTNNRQTESQIMSEFP
						FTIATKRIKYLGIQLTRDVKDLF
		l				KENYKPLLKEIREDTNKWKNIP
						CSRIGRINIMKMAILPKVIYRFN
						DIPIKLPMTFFTELEKTTLKFIW
						NQKRACIAKTILSKKNIAGGITL
						PDFKLYYKATVTKTAWYWYQ
						NRDIDQWNRTEASEVTSHIYNH
	}					LIFYKPDKNKKWGNDSLFNKW
		:				CWENWLAICRKLKLDPFLTPYT
						KIHSRWIKDLNVRPKTIKTLEEN
						LGNTIQDIGMGKDFMTKTPKA
						MATKAKVDKWDVIKLKSFCTA
		1				KETTIRVSRQPTEWEKIFAIYPS
						DKGLISRIYKELKQIYRKK\TNN
		1				PIKKWAKNMNRHFSKEDIYAA
						NRQMKKCSSSLVIREMQIKTTM

SEQ ID NO:	SEQ ID NO: of peptide	Met	SEQ ID NO: in USSN	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
5617	35985	l la	5663	1391	2742	KKRESSLTHVMRPASF*YQSQA
3017	33963		3003	1371	2142	ETQQKKRILDQYP**TLMQKSSI
						K/YLAKRIQQHIKKLIHHDQVGF
:						IPGMQGWFNIRKSINVIQHINRA
		l				KDKNHMIISIDAEKAFDKIQQPF
•						MLKTLNKLELEETTVKFIWNQ
						KRAHIAKSILSQKNKAGGITLR
						DFKLYYKATVTKTAWYWYQN
						RDIDQWNRTEPSEITLHIYNYLI
						FDKPEKNKQWGTDSLFNKWC
			•			WENWLAICRKLKLDPFLTPYTK
						INSRWIKDLNVRPKTIKTLEENL
						GITIQDIGMGKDFMSKTPKAMA
						TKAKIDKWDLIKLKSFCTAKET
						TIRVNRQPTKWEKIFTTYSSDK
						GLISRIYNELKQIYKKKTNNPIK
		ŀ				KWAKDMNRHFSEEDIYAAKKH
						MKKCSSSLAIREMQIKTTMRYH
						LTPVRMAIIKKSGNNRHAPFSIH
						THIMFGSLYLQIQKDLSILGFW
		<u> </u>				YPRGILEPIY
5618	35986	Α	5664	1	1910	MTIIINIIIIIIIIIIIIIIVHIITINIVIT
						TTILTTVIISSTIIIITIFVFETGAI
						KLEIRIKKLTQNRSATWKLNNL
						LLNDYWVHNEMKAEIKMFFET
						NENKOTTCQNLWDTFKAINKID
						RPLARLIKKKREKNQIDAIKND
	ļ					KGDITTDTTGIQTTIREYYKHLY
						ANKLENREEMDKFFDTYTLPRL NQEEVESLNRPITGSKIEEIINSL
						SIKKSPGPDGFTAEFYQRYKEE
						LVPFLLKLFQSIEKEGIILNSCCE
						ASIILIPKPGRDTTKKENFRPISL
						MNIDAKILNKTLANRIQQHITK
						LIHHVQVGFIPGMQGWFNIRKS
						INNWKKTTLKLIWNQKRACITK
						SILSQKNKAGGITLPDFKLYYK
						ATVTKTAWYWYQNRDIDQWN
						RTEASEIIPHTYNYLSFDKPDKN
						KKWGNNSLFNKWCWENWLAI
						CRKLKLDPFLTLYTKINSRWIK
						DLNFGTETIKTLEENLGNTIHDI
						GMGKDFMSKTPKAKATKAKID
						KWDLIKLKSFCTAKETTIRVNR
						QPTEWEKIFAIYPSDKGLISRIY
ŀ						KELKQVYKKKTNNPIKKWAKD
						TNRHFSNEDIYAANRHMKKCS
						SSLAIREMQIKTTMRYHLAPVR
						MTIIKKSGSNRQ/W*ASGHVRC

NO: of peptide sequence hod in USSN docation of first codon for last amino acid feetion, impossible nu sequence seque	(X=Unknown,
5619 35987 A 5665 I 2460 MPHAGHARSPGI MSRWMEWNPGI DLPRDIQFDSEKG ALNQEEVESLNR SLPTQKSSGPDGI EELVPFLLKLFQG YEASIILIPKPGR SLMNIDAKILNK KLIHQDQVGIIPC SINVIQHINRTKE KAFDKIQQPFML YLRIQLTRDVKD NEIKEDTINKWKI IIKMA/LPKVIVR FFSELEKSTLKFI TILSQKNKAGGII ATVTKTAWYWY RTEPSEMTPHIY KQWGKDSLFNK GRQLKLDPFLTP DLNVRPKTIKTLL SMGKDPFKSKTPI DLNVRPKTIKTLL SMGKDPKSKTPI DKWDLIKLKSFC RQPTEWEKNFAJ YKQLKQIYKKKI DMNRHFSKEDV CSSSLAIREMQIK TMAJIKKSGNNR TLLYCWWDCKL QELRDLEGIPFE DYKSCCYKDTC" HILKQHRRKELE DYKSCCYKDTC" HILKQHRRKELE LEENLGNTUR KWLLVKMAIL/PQAIC TYTKINSRWIKD LEENLGNTIQDIG PKAMATEAKIDK	
MSRWMEWNPGI DLPRDIQFDSEK ALNQEEVESLNR SLPTQKSSGPDGI EELVPFLLKLFQI YEASIILIPKPGR' SLMNIDAKILINK KLIHQDQVGIIPC SINVIQHINRITKE KAFDKIQQPFML YLRIQLTRDVKDI NEIKEDTINKWKI IIKMAI/LPKVIYF FFSELEKSTLKFI TILSQKNKAGGII ATVTKTAWYWY RTEPSEMTPHIYY KQWGKDSJENK GRQLKLDPFLTP DLNVRPKTIKTL SMGKDFMSKTPI DKWDLIKLKSFC RQPTEWEKNFAI YKQLKQIYKKKI DMNRHFSKEDV' CSSSLAIREMQIK TMAIIKKSGNNR TLLYCWWDCKL QFLRDLELGIPFE DYKSCCYKDTC' HILKQHRRKELE 5620 35988 A 5666 689 1909 LIAYQPKKSRTR KYLGIQLTRDVK LNEIKEDINKWK LVKMAIL/PQAIC TYTKINSRWIKD LEENLGNTIQDIG TYTKINSRWIKD LEENLGNTIQDIG TYTKINSRWIKD LEENLGNTIQDIG PKAMATEAKIDK	,
MSRWMEWNPGI DLPRDIQFDSEK ALNQEEVESLNR SLPTQKSSGPDGI EELVPFLLKLFQI YEASIILIPKPGR' SLMNIDAKILINK KLIHQDQVGIIPC SINVIQHINRITKE KAFDKIQQPFML YLRIQLTRDVKDI NEIKEDTINKWKI IIKMAI/LPKVIYF FFSELEKSTLKFI TILSQKNKAGGII ATVTKTAWYWY RTEPSEMTPHIYY KQWGKDSJENK GRQLKLDPFLTP DLNVRPKTIKTL SMGKDFMSKTPI DKWDLIKLKSFC RQPTEWEKNFAI YKQLKQIYKKKI DMNRHFSKEDV' CSSSLAIREMQIK TMAIIKKSGNNR TLLYCWWDCKL QFLRDLELGIPFE DYKSCCYKDTC' HILKQHRRKELE 5620 35988 A 5666 689 1909 LIAYQPKKSRTR KYLGIQLTRDVK LNEIKEDINKWK LVKMAIL/PQAIC TYTKINSRWIKD LEENLGNTIQDIG TYTKINSRWIKD LEENLGNTIQDIG TYTKINSRWIKD LEENLGNTIQDIG PKAMATEAKIDK	HTUKLACWW
DLPRDIQFDSEKG ALNQEEVESLNR SLPTQKSSGPPG EELVPFLLKLFQ FEASIILIPKPGRN SLMNIDAKILNK KLIHQDQVGIIIPK SINVIQHINQPVMIL YLRIQLTRDVKD NEIKEDTINKWKI IIKMAI/LPKVIYF FFSELEKSTLKFI TILSQKNKAGGII ATVTKTA WWW RTEPSEMTPHIY KQWGKDSLFNK GRQKLDFFLTP DLNVRPKTIKTL SMGKDFMSKTPI DK WDLIKLKSFO RQPTEWEKNFAI YKQLKQIYKKKI DMNRHFSKEDV CSSSLAIREMQIK TMAIIKKSGNR TLLYCWDCKL QFLRDLELGIPFE DYKSCCYKDTC' HILKQHRRKEI 5620 35988 A 5666 689 1909 LIAYQPKSRTR KYLGIQLTRDVK LNEIKEDINKWK LVKMAIL/PQAIC TYTKINSRWIKK LEENLGNTIQDIG PKAMATEAKIDK	
ALNQEEVESLNR SLPTQKSSGPDG EELVPFLLKLFGY YEASIILPKPGRY SLMNIDAKILNK KLIHQDQVGIIPC SINVIQHINRTKE KAFDKIQQPFWL YLRIQLTRDVKD NEIKEBTINKWKY IIKMAI/LPKVIYR FFSELEKSTLKFI TILSQKNKAGGI ATVTKTA WYWY RTEPSEMTPHIYY KQWGKDSLFNK GRQLKLDPFLTP DLNVRPKTIKTL SMGKDFMSKTPI DK WDLIKLKSFC RQPTEWEKNFAI YKQLKQIYKKKY DMNRHFSKEDD CSSSLAIREMQIK TMAIIKKSGNNR TLLYCWWDCKL QFLRDLELGIPFE DYKSCCYKDTE HILKQHRRKELE 5620 35988 A 5666 689 1909 LIAYQPKKSSTR KYLGIQLTRDVK LNEIKEDINKWK LVKMAIL/PQAIC TYTKINSRWIKD LEENLGNTIQDIC PKAMATEAKIDK	
SLPTQKSSGPDGI EELVPFLLKLFQS YEASIILIPKPGS SLMNIDAKILNK KLIHQDQVGIIPC SINVIQHINRTKE KAFDKIQQPFML YLRIQLTRDVKM NEIKEDTNKWKN IIKMAI/LPKVIYR FFSELEKSTLKFI TILSQKNKAGGII ATVTKTA WYW RTEPSEMTPHIYY KQWGKDSLFNK GRQLKLDPFLTP DLNVRPKTIKTL SMGKDFMSKTPI DK WDLIKLKSFC RQPTEWEKNFAI YKQLKQIYKKK: DMNRHFSKEDD CSSSLAIREMQIK TMAIIKKSGNNR TLLYCWWDCKL QFLRDLELGIPFE DYKSCCYKDTC HILKQHRRKELE HILKQHRRKELE LIAYQPKKSRTR KYLGIQLTRDVK LNEIKEDINKWK LVKMAIL/PQAIC TYTKINSRWIKD LEENLGNTIQDIC PKAMATEAKIDK	
EELVPFLLKLFQE YEASIILIPKPGRS SLMNIDAKILNK KLIHQDQVGIIPC SINVIQHINRTKDE KAFDKIQQPFML YLRIQLTRDVKD NEIKEDTNKWKS IIKMAI/LPKVIYR FFFSELEKSTLKFT! TILSQKNKAGGII ATVTKTAWYWS RTEPSEMTPHIYS KQWGKDSLFNK GRQLKLDPFLTP DLNVRPKTIKTL. SMGKDFMSKTPI DK WDLIKLKSFC RQPTEWEKNFAI YKQLKQIYKKKS DDMNRHFSKEDV CSSSLAIREMQIK TMAIIKKSGNNR TLLYCWWDCKL QFLRDLELGIIPFE DYKSCCYKDTCS HILKQHRRKELE 5620 35988 A 5666 689 1909 LIAYQPKKSRTR KYLGIQLTRDVK LNEIKEDINKWK LVKMAILPQAIC TYTKINSRWIKD LEENLGNTIQDIC TYTKINSRWIKD LEENLGNTIQDIC TYTKINSRWIKD LEENLGNTIQDIC	
YEASIILIPKPGRY SLMNIDAKILNK KLIHQDQVGIPC SINVIQHINRTKE KAFDKIQQPFML YLRQLTRDVKD NEIKEDTNKWKI IIKMAI/LPKVIYR FFSELEKSTLKFI TILSQKNKAGGII ATVTKTAWYWY RTEPSEMTPHIYY KQWGKDSLFNK GRQLKLDPFLTP DLNVRPKTIKTL SMGKDFNSKTPI DK WDLIKLKSFG RQPTEWEKNFAI YKQLKQIYKKKI DMNRHFSKEDV CSSSLAIREMQIK TMAIIKKSGNNR TLLYCWWDCKL QFLRDLELGIPPE DYKSCCYKDTC* HILKQHRRKELE 5620 35988 A 5666 689 1909 LIAYQPKKSTR' KYLGIQLTRDVK LNEIKEDINKWK LVKMAIL/PQAIC TYYTKINSRWIKD LEENLGNTIQDIC PKAMATEAKIDK	•
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SMGKDFMSKTPI DK WDLIKLKSFO RQPTEWEKNFAI YKQLKQIYKKKT DMNRHFSKEDV CSSSLAIREMQIK TMAIIKKSGNNR TLLYCWWDCKL QFLRDLELGIPFE DYKSCCYKDTCT HILKQHRRKELE 5620 35988 A 5666 689 1909 LIAYQPKKSRTR KYLGIQLTRDVK LNEIKEDINKWK LVKMAIL/PQAIC TYTKINSRWIKD LEENLGNTIQDIC PKAMATEAKIDK	
DKWDLIKLKSFO RQPTEWEKNFAI YKQLKQIYKKKT DMNRHFSKEDV CSSSLAIREMQIK TMAIIKKSGNNR TLLYCWWDCKL QFLRDLELGIPFE DYKSCCYKDTCT HILKQHRRKELE 5620 35988 A 5666 689 1909 LIAYQPKKSRTR KYLGIQLTRDVK LNEIKEDINKWK LVKMAIL/PQAIC TYTKINSRWIKD LEENLGNTIQDIC PKAMATEAKIDK	-
RQPTEWEKNFAI YKQLKQIYKKKT DMNRHFSKEDV CSSSLAIREMQIK TMAIIKKSGNNR TLLYCWWDCKL QFLRDLELGIPFE DYKSCCYKDTCT HILKQHRRKELE 5620 35988 A 5666 689 1909 LIAYQPKKSRTR KYLGIQLTRDVK LNEIKEDINKWK LVKMAIL/PQAIC TYTKINSRWIKD LEENLGNTIQDIC PKAMATEAKIDK	
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DYKSCCYKDTCT HILKQHRRKELE 5620 35988 A 5666 689 1909 LIAYQPKKSRTR KYLGIQLTRDVK LNEIKEDINKWK LVKMAIL/PQAIC TYTKINSRWIKD LEENLGNTIQDIC PKAMATEAKIDK	VQPLWKTLW
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5620 35988 A 5666 689 1909 LIAYQPKKSRTR' KYLGIQLTRDVK LNEIKEDINKWK LVKMAIL/PQAIC TYTKINSRWIKD LEENLGNTIQDIC PKAMATEAKIDK	TPKLARDDQI
KYLGIQLTRDVK LNEIKEDINKWK LVKMAIL/PQAIC TYTKINSRWIKD LEENLGNTIQDIC PKAMATEAKIDK	TRQKQYRAW
LNEIKEDINKWK LVKMAIL/PQAIC TYTKINSRWIKD LEENLGNTIQDIC PKAMATEAKIDK	•
LVKMAIL/PQAIC TYTKINSRWIKD LEENLGNTIQDIC PKAMATEAKIDK	
TYTKINSRWIKD LEENLGNTIQDIC PKAMATEAKIDK	
LEENLGNTIQDIC PKAMATEAKIDK	
PKAMATEAKIDK	
AAKETTTRVNR(
YAANRHMKKCS	
TTVSPFAAQWVS	•
HPSNNDGKGYRO	•
VHGPQDQKKIIK	-
NILMNAKTIENG	
QQPHDAENLFT	-
SPCIEVSAALSQV	
GIPNRILRSPIPGL	
VWVRGDPTALK	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5621	35989	В	5667	1	1962	
5622	35990	В	5668	1	1851	
5623	35991	A	5669		1947	MNIDAKILNKILANRIQQHIKKL IHHDQVGFIPGRQGWFNICKSIN VIQHINRAKDKNHMIISIDAEKA FDKIQQLFMLKTLNKLGIDGTY FKIIRAIYDKPTANIILNGKKLEA FPLKTGTRQGCPLSPLLFNIVLE VLARAIRQEKEIKGIQLGKEEV KLSLFADDMIVYLENPIVSAQN LLKLISNFSKVSGYKINVQKSQ AFLYTNNRQTESQIMSELPFTIA SKRIKYLGIQLTRDVKDLFKEN YKPLLKEIKEDANKWKNIPCSW VGRINIVKMAILPKVIYRFNAIPI KLPMTFFTELEKTTLKFIWNQK RAHITKAILSQKNKARGITLPDF KLYYKATVTKTAWYWYQNRD IDQWNRTQPSEITPHIYNYLIFD
5624	35000		5670		1067	KPDKNKQWGKGSLFNKWCWE NWLAICRKLKLDPFLTPYTKIN SRWIKDLNVRPKTTKTLEENLG ITIQDIGMGMDFMSKTPKAMAT KDKIDKWDLIKLKSFCTAKETT IRVNRQPTKWEKIFTTYSSDKG LISRIYNELKQIYKKKTNNPIKK WAKDMNRHFSKEDIYAAKKH MKKCSPSLAIREMQIKTTMRYH LTPVRMAIIKKSGNNRCWRGC GEIGTLLHCW\WINWMKKTWH IYTMEYYASIKKNEFMSFAGA* MKLETII
5624	35992	Α	5670	2	1967	

SEQ ID NO:	of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5625	35993	A	5671	1039	2272	KKRESSLTHVMRPASF*YQSQA ETQQKKRILDQYP**TLMQKSSI K/YLAKRIQQHIKKLIHHDQVGF IPGMQGWFNIRKSINVIQHINRA KDKNHMIISIDAEKAFDKIQQPF MLKTLNKLELEETTVKFIWNQ KRAHIAKSILSQKNKAGGITLR DFKLYYKATVTKTAWYWYQN RDIDQWNRTEPSEITLHIYNYLI FDKPEKNKQWGTDSLFNKWC WENWLAICRKLKLDPFLTPYTK INSRWIKDLNVRPKTIKTLEENL GITIQDIGMGKDFMSKTPKAMA TKAKIDKWDLIKLKSFCTAKET TIRVNRQPTKWEKIFTTYSSDK GLISRIYNELKQIYKKKTNNPIK KWAKDMNRHFSEEDIYAAKKH MKKCSSSLAIREMQIKTTMRYH LTPVRMAIIKKSGNN
5626	35994	Α	5672	1	3477	
5627	35995	Α	5673	1	2814	
5628	35996	Α	5674	1	2093	
5629	35997	Α	5675	1	2724	
5630	35998	Α	5676		4680	MDKFLDTYTLPRLNQEEVESLN RPITGSEIVAIINSLPTKKDPGPD GFTAEFYQRIKYLGIQLTRDVK DLFKENYKPLLKEIKEDTNKW KNIPCSWVGRINIMKMAILPKVI YRFNAIPIKLPMTFFTELEKTTL KFIWNQKRARITKSILSQKNKA GGITLPDFKLYYKATVTKTAW YWYQNRDIDQWNRAEPSEIML PIYNYLIFDKPDKNKQWGKDSL FNKWCWENWLAICRKLKLDPF LTPYTKINSRWIK
5631	35999	В	5677	1	1989	
5632	36000	Α	5678	1	2784	
5633	36001	В	5679	1	3573	
5634	36002	В	5680	1	2630	
5635	36003	Α	5681	5339	10577	
5636	36004	Α	5682	1	873	
5637	36005	Α	5683	1	672	
5638	36006	A	5684	1	435	

SEQ ID			SEQ ID NO:			Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
		ŀ		sequence		
5639	36007	Α	5685	1	1011	MVRFGDELGGRYGGPGGERA
						RGGGAGGAGGPGPGGLQPGQR
						VLYKQSIAQRARTMALYNPIPV
		ŀ				KQNCFTVNRSLFVFSEDNVVRK
						YAKRITEWPPFEYMILATIIANC
						IVLALEQHLPDGDKTPMSERLD
						DTEPYFIGIFCFEAGIKIIALGFV
		l				FHKGSYLRNGWNVMDFVVVL
						TGRKAGLGCCGSGVESGGWGD
						RSSSPAMEQSRMENDFDELTEV
		1				GFRKSVITNFSELKE\DVRTHRK
						EAKNLEKRLDKM\VNRINSVEK
		l				TLNDPMELKTMARELCDACTS
						FSS*FNQVEEKVSVIEDQMDEM
		l				K*EEKFREKRVKRNEQSLQEIW
		l				DYVKRPNLRL
5640	36008	A	5686	1	1539	
5641	36009	Α	5687	1	756	MDDPRLNVKPLTESLETYSGKS
						KGLEGDVITDSKLNDCRCPSGT
		l				KLPEEGLGSNICCSAIFAVLQPL
		l				LVIPRQTGSGVDHQQTPTDLRM
						TVRRKMNKQKGIASTSTKRTK
		1				KDIHTKTPSVCHQHQRPKDCSP
						SPATEQSWMENDLEELTEIGFR
						RSVITNFSELKEDVRIHHKKAK
		l				NLGKRLDEWLTRINSVEKTLND
						LMELKTMARELHDSCTSFNSRS
						NQAEEKVSVIEDQFNEIKREEK
İ		1				FREK\RVKRN
5642	36010	Α	5688	1	1008	MVRFGDELGGRYGGPGGERA
						RGGGAGGAGGPGPGGLQPGQR
						VLYKQSIAQRARTMALYNPIPV
		•				KQNCFTVNRSLFVFSEDNVVRK
						YAKRITEWPPFEYMILATHANC
						IVLALEQHLPDGDKTPMSERLD
	1	1				DTEPYFIGIFCFEAGIKIIALGFV
						FHKGSYLRNGWNVMDFVVVL
		l			1	TGRKAGLGCCGSGVESGGWGD
						RSSSPAMEQSRMENDFDELTEV
				,		GFRKSVITNFSELKEDVRTHRK
						EAKNLEKRLDEWLTRMNSVEK
						TLNDLMELKTMARELRDACTS
1						FSSQFDQVQEMVSVIQDQISEM
						KREEKFREKRVKRNE/QSLQ*I
						WDYVERPNLRL
5643	36011	A	5689	1	672	
5644	36012	A	5690	1	807	
5645	36013	Α	5691	1	1617	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5646	36014	A	5692	1064	1893	LLEGSININKKDIYTKTQSVGRQ HQRPKVDKTTKMGRNQNRKA ENSKNQSASSPPKDC\SSSPAME QSWMENNFDELTEIGFRRSVIT NFSELKEDVQTHCKEAKNLKK RLDECLTRINSVEKKTLNDLME LKTMAQELHDTCTSFNTQFDQ VEERVSVIEDQINEIK*EEKFRE K\RVKINE*SCQEIWD*VKRPNI HLITVPESDGEKGTKLENTLQDI IQENFHNIARQANIQIQEIQRTA QRYSSRTATPRHIIVRFAKVEM KEKVLRAARKKG
5647	36015	A	5693	3	620	EAGWLLLSSVDEVMKENDELR DSISQLQKQILSLKSAKIALTESL ISFRERAEIVEKQTQALIM*VAD LQQGNDQTFQGLLDTGSELTLI PGDPKHHCGPPGPKVRAYGDQ VINGVLAQVQLIVGPVGPWTHP VVISPVPECVIGIAILNSWQNPHI GSLTGRVRAIMVGISWQGQQY TFTVLPQRYINFPALCHKTAKR HTARR
5648	36016	Α	5694	1	1098	
5649	36017	٨	5695	2	563	IIRRAVFRWFWG*RSWGLNMIR PQNTKDSNSNCLEKTDSPWCEL FKELYKINAFDTPDSLLMRGNE FSDPIHHTFDHMWRTKEHNEA GWLLLSSVDKVMKENDELGDS ISQLQKQILSLKSAKIALSESLIS CRERAEIVEK*T*ALIMLVADL Q*KVHAQPHHAQPRQVSTVKV RALIVLQEVTDP
5650	36018	A	5696	I	499	FVLVDVNNMWHRYASMLYER RLLIICSKLSTLTACIHGFAAML YPMYWQHVYI\PVLPPHLLDYC CAPMPYLIGIHLSLMEKVRNMA LDDVVILNVDTNTLETPFDDLQ SLPNDVISSLKNRLKKVSTTTG DGVARAFLKAQAAFFGSYRNA LKIEPEEPITFCEEA
5651	36019	Α	5697	1	795	
5652	36020	A	5698	103	3531	
5653	36021	A	5699	1	2073	
5654	36022	A	5700	140	901	
5655	36023 36024	A	5701	1	639 318	REYGTPQEQLNLALLTLNFLSL PKGQMLSAAEQHLQKPAAKTE A/GRMIWQRDPITKIWEIGKIIT WGRGYACVSPGQNHQSVWIPS RHLKSCHGPDAKEEIPGGS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
		ļ		sequence		
5657	36025	I A	5703	<u> </u>	612	PTRPIKHITDIPYNSEGQAIVER
3037	30023	^	3703		012	MNLSLKQQLQKHKKGGK/SKD
						YGTPHMQLNLALLTLNFLSLPK
						GQILSVAEQHLQKPAAKTEAEQ
						LVWWRDPITKS*EIGKIITWGK
	1					·
						GYACVSPGLTQQPIWIPSRHLK
						PYHEPDAEEEIPGGSRGPPPPPV/
	İ	1				DSHVETDAEEEPHCHEQHLSNT
	ı	1				ATHLGTDQEAVTAGGRKPEES
	2.026	<u> </u>	570.4		701	KTTSHNK
5658	36026	Α	5704	2	791	DCCOLLAL LIBLAL DCL COACH
5659	36027	Α	5705	2301	2620	DGQQLIALHRLALRELQQAGH
				ŀ		AGLPQQAKILFDGGSEIGKIRGL
						QRPRAKNRLSGRGPLREPSPK*F
						FGVQGRKTLNRTLKGAPH*NL
		ļ				AAKAGKPLSPCTSGRIRM
5660	36028	Α	5706]1	1010	MDDIPQEAGRYRHNQAYAYSI
						QGDGAEDDDERIVRFHTRVTV
						DSDTLASDAARLTCRHGLGNQ
			İ			GSDLVLYGTSTPKNLIWVRVH
						VVGHRPDRRFFAFDVWSPRSLI
						VDSCSKLEQHSTLSRAILLIYKG
						FCRFRNHHQTGFSPAGANQRGP
						LAATLSGPGGEGQSAVARLTGE
	1					KKNHPGAQYANRLSPRVGRFIN
						AAGTTGFPTGKRAVSATQLMD
						FADFGTTIKQDFRLLGQTSVDR
						LLQLSQGQANVQQFIDEGNYTS
	1					GDNHTLRDPHYVEDKGHKYLV
		İ			<u> </u>	FEANTGTENGYQGEESLFNKA
						YYGGGTNFFRKESQKLQQSAK
	1					KRDAELANGALGIIELNNDYTL
	-					KKVMKPLITSNTVTDEIERANV
						FKMNGKWYLFTDSRGSKMTID
				•		GINSNDIYMLGYVSNSLTGPYK
						PLNKTGLVLQMGLDPNDVTFT
						YSHFAVPQAKGNNVVITSYMT
				,		RGSSR*RQRP*IPCIRSQHGNRK
						RIPR/SKNLYLTKRTTAAARTSS
						VKKARSFSRALKNAMLS*RTAP
5661	36029	В	5707	1	1149	
5662	36030	В	5708	55	3947	
5663	36031	В	5709	4	885	
	30031	12	3703	1		<u> </u>

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
5664	36032	A	5710	461	1560	SLQTRDLPISEPPSNRIFACWGK
	1					PAWTACCNSLRARR*RAISCCP
						SH*KRIPRRRIFI*PKRTTAAART
						SSVKKARSFSRALKNAMLS*RT
						APSVS*S*IMITH*KK**SR*SLQ
						TRDLPISEPPSNR\FSPAGANQR
						GPLAATLSGPGGEGQSAVARLT
•			}			GEKKNHPGAQYANRLSPRVGR
[1					FINAAGTTGIPDWKAGSERNAI
						NGAIALIEGFRPLTLEPSSLIVNS
		İ				CSKLEQHSTLSRAILLIYKGFCR
						FRNHHQTGFSPAGPNQRGPLAA
						•
						TLSGPGGEGQSAVARLTGEKK
						NHPGAQYANRLSPRVGRFINAA
	1	<u> </u>		<u> </u>	1450	GTTGFPTGKRAVSATQLM
5665	36033	В	5711	1	1458	
5666	36034	В	5712	1	3573	0.07.0.000.000.000.7777.0
5667	36035	Α	5713	606	1005	GGMLCGTLSGPDGEGQSAVAR
						LTGEKKHPRGDQ*QKAPLLAR
						W/RNPLMQLAQ/PDFPTGKRAI
						DTHRSVSHAHIIITIHSSTLSSDIS
						HTDAPNTSSNNYVIAITSPSDTA
						PTSRTPLHRIHHTTPIPIATLARP
5668	36036	В	5714	1	1566	
5669	36037	С	5715	29	2314	
5670	36038	Α	5716	597	1384	ISYTVPEALSARSNPAACQHAS
						CRPAACPVASVGSPRQPAPGGA
						GGHRHWPRRRWRLRQRPRPSA
ļ		:				PGRPRRPRPRPRRHRRRPRPHP
						HHAPPPRPSCAETRS\RRGGGRR
		•				PLPRPLPPGAAGAGHREVGDK
		ŀ				AGAAGCGRRVVALAA\PVSPPS
					İ	SPSSPSPSLLSAAVGVAGCFSAE
						LSDSRRLLAPGGEGPLARSPAR
						CDPAAGSACPAGWSSRQRSRC
						ELMCAAGISCWHTGHCTWCSL
						SFMPLVLMILAASFRPVAVSSH
5671	36039	В	5717	1	3873	
5672	36040	С	5718	356	784	
5673	36041	Α	5719	1303	1664	KRRDEIIPCVLSKQLLVTFITCR
						ASCINESANARGEAVCVLGAR
						VV/SSFHQ*DGQQLIALHRLALR
1						ELQQAVHAGLPQQAKILFDGGS
						EIGKIPYKSKE*PENRLSVVPVW
						EQESTLASV
						EQESTLASV

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5674	36042	A	5720	453	915	TVSRWAMLEHILLFSREPSNKE PET*AHSM*ADDKRFAAYKKK TKGGTERWDKYGRRPANGIRT
						GENNAKTEHLRDRDQKR/PRER ERAR*TTEITEQEGSG*GEPVEQ MRKRRAQAWSRRKSTGSQTET RSVRVTRVGLKTASESRRIRTM EQN
5675	36043	В	5721	319	1469	17211
5676	36044	В	5722	164	3112	
5677	36045	A	5723		1102	MRWTPGHEITKALQTFQRYTGI QHVHRIGMAERMWCDRNRER HTVSSSGGNRLPNPEHFEAFQS VAQCTYNQTVQLDITAFLKTV KKNKHKFYPAFIHILARLMNAH PEFRMAMKDVDSCSKLEQHST LSRAILLIYKGFCRFRNHHQTGF SPAGANQRGPLAATLSGPGGEG QSAVARLTGEKKNHPGAQYAN RLSPRVGRFINAAGTTGFPTGK RAVSATQLIYRELFPQPLVQLD PCSVPCMQPPTSPKSRAWVSDI ALLEFQKKKEMEKKKEKERKK ELQGGRGRRRGGGWGKGRER GGRGKKRRGGGEAIRDAEKAG RLPHPDMEIRGRVEQRVGYTIE QINHMRDVFGTRLRRAEDVFPP VIGVAAHKGGVYKTSVSVHLA QDLALKGLRVLLVEGNDPQGT ASMYHGWVPDLHIHAEDTLLP FYLGEKDDVTYAIKPTCWPGL DIIPSCLALHRIETELMGKFDEG KLPTDPHLMLRLAIETVAHDYD VIVIDSAPNLGIGTINVHPAWGS AISLLEYSPGHLAGFVASIPAPL QLVPRWDTKPVLEDASASSRFS IPQRNPTHLHPTTSFPSCNPDTV DAWMRFLITWCLYTSPYSTLM GELFPFYQGMLLESTKQGSQVR KVFVWTLVPNWNNTQPYSRAI LLIYKGFCRFRNHHQTGFSPAG
5670	26046	D	5724	90	6258	ANQRGPLAATLSGPGGEGQSA
5678 5679	36046 36047	B A	5724 5725	837	1485	
5680	36047	A	5726	1	2993	
5681	36049	A	5727	1	123	
5682	36050	A	5728	1	960	
5683	36051	A	5729	507	755	NKRGYKQMEEHSMLMGRKNQ YRKNGHTAQGNL*IQCHPHQA TNDFLHRIGKNDFQVHMEPKK SPHRQVNPKPKEQSWRHHAT

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5684	36052	A	5730	274	882	PDQVDRGGAHWQTDADDPWL ADHLQHCQRCGEILRHYSGGIR GNVSAVKCAEHHVPAFARLRN PQCGDFQRLDYRLFDSPGVKRR EL*TAYRFCHVAP*LMDLRSG WAAGAVYRYQSH*FTADRLRS\ VDMDEAGNHHSQQT\SQDQKI KQRTFSLISGS*TTRTHGHREGN ITHRGLSRPNTARILCASKAHLG LPLRSSLP
5685	36053	Α	5731	114	329	
5686	36054	С	5732	1	1983	
5687	36055	C	5733	1	588	
5688	36056	Α	5734	1	597	
5689	36057	Α	5735		721	MTSYNEQNKKPVTDPNEMAIH EDSNQEFKIAVVNETQRSPRSSP TEHLMTPPPLPTREQQPPLTVIF HYLPKSYKTAPPLSPFPDSLFGL SQPAPRDMDEAGNHHSQQT\K QEQKTKHCMFSLTSGS*TMRTH GHREGTVIHRGLLGQFSQQLIN KSEFSNTVCREKSGRVLQVNM ESPL*MIQDALQDQTTPKGPITV FL*LFYLDHLLRSSHKVASYLST ESRSRGSITVLFIRCSEP
5690	36058	A	5736	417	696	DRNRKANTACFQHSQVGVEQ* EHMDTGKV*SI*YGNYEHTSTC ERMEIQKATTEHLNL*S*SGSFL RFTWKMLQFSKISRSFYQVLKH QQKIM
5691	36059	A	5737	325	1489	RSPTLLMHVHYIKVISWRELLLI WLRAANASINPGVEQSPSGDPK TVTGFPFLRLEAADWVEEASVP KGVHIALSSQEQRAGDPRDLEA PSNLVISERTHRSFRVSWTPPSD SVDRYKVEYYPVSGGKRQEVV RTPGVPDLRLPAIIPQAAWLLA HLPAPSAPPTELPTKHTEAQSQ RCWSLLLCSVTEVAAPSQVVK EAPPSFHTGTENGYQGEESLFN KAYYGGGTNFFRKESQKLQQS AKKRDAELANGALGIIELNNDY TLKKVMKPLITSNTVTDEIERA NVFKMNGKWYLFTDSRGSKM TIDENKIPRNPTYKGREGPLQGE LQTTAQGNKRGYKQMEEHSM LMDRKNQYRENGLTAQA\PQPP PSGI*GYTAPCFPWENLI
5692	36060	В	5738	1	873	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
5693	36061	A	5739	274	716	QACYIYTTEYYAAIKNDEF/TVL
						CRDMDEAGNHHSQQTVTRTKN
		1		ļ		QTPHVLTHRWELNNENTWTQE
	Ì					GEHHTPGPVMLDSAPPTLGHQT
						PGSSAFGLWDLHQRLRLPRPQT
						KGSTVGFPGSEAFKLGLGHYW
		1				LSFFPSLQTAYRGTLPFNH
5694	36062	В	5740	134	2283	
5695	36063	Α	5741	3	872	
5696	36064	Α	5742	1	480	SKLNSGDGCWSGARRGSAHSR
,						SPLPSPLRPRAKMATEGLHENE
						TLASLKSEAESLKGKLEEERAK
						LHDVELHQVAERV/VGPGAVC
		İ				HEDQKDPQRPREQSPVHGLVQ
						R*EEDRELVTGWEGDRVGFLH
		1		}		HKQGARGHHALHVGDGMCLC
				ŀ		PIGMCHCLWWFG
5697	36065	A	5743	3	387	AKMATEGLHENETLASLKSEA
						ESLKGKLEEERAKLHDVELHQ
						VAERV/VGPGAVCHEDQKDPQ
				-		RPREQSPVHGLVQR*EEDRELV
						TGWEGDRVGFLHHKQDPDSER
						RWHMCPVGRGERAAAAELPRT
5698	36066	Α	5744	1	1575	
5699	36067	Α	5745	20	510	
5700	36068	Α	5746	1	1392	
5701	36069	Α	5747	1	1560	
5702	36070	Α	5748	3	325	
5703	36071	С	5749	206	409	
5704	36072	Α	5750	1	855	
5705	36073	С	5751	233	571	
5706	36074	A	5752	1	3600	
5707	36075	Α	5753	l .	4137	
5708	36076	Α	5754	3	1357	
5709	36077	A	5755	3	194	•
5710	36078	Α	5756	3	269	
5711	36079	A	5757	1	1770	
5712	36080	Α	5758	164	411	RKNRKPNTAFLTHKWELNNEN
						TWTQGGNITHQG\CWGVQG*R
						RDSIRRNT*CKCRVPGCSKPPW
						HVYTYVTNLHILHMYPRT

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5713	36081	Α	5759	1	3095	MKMADAKQKRNEQLKRWIGS
						ETDLEPPVVKRQKTKVKFDDG
	i	l				AVFLAACSSGDTDEVLKLLHR
						GADINYANVDGLTALHQACID
		l				DNVDMVKFLVENGANINQPDN
		ı				EGWIPLHAAASCGYLDIAEFLIG
				:		QGAHVGAVNSEGDTPLDIAEEE
						AMEELLQNEVNRQGVDIEAAR
						KEEERIMLRDARQWLNSGHIN
:		ļ				DVRHAKSGGTALHVAAAKGY
ľ						TEVLKLLIQAGYDVNIKDYDG
						WTPLHAAAHWGKEEACRILVD
5714	36082	Α	5760	1	591	FTVTRCYAVAQEGTYFDGSGY
		1				AALVKEGYKVQSDVNITLEVS
						N\PRRNGVFLGISTAIVHAIGLEL
						VDGKVLFHVNNGAGRITAAYE
		ł				PKTATVLCDGKWHTLQANKSK
		İ				HRITLIVDGNAVGAESPHTQSTS
		1				VDTNNPIYVGGYPAGVKQKCL
				}		RSQTSFRGCLRKLALIKSPQVQS
!		ł				FDFSRAFELHGVFLHSCPGTES
5715	36083	Α	5761	340	739	
5716	36084	A	5762	1234	1897	
5717	36085	A	5763	1	4070	VTPRAAWLGLGFRGSAVLGLC
	ľ					WQPRSPPSRAAGMMNRTTPDQ
						ELVPASEPVWERPWSVEEIRRS
						SQSWSLAADAGLLQFLQEFSQQ
						TISRTHEIKKQVDGLIRETKATD
						CRLHNVFNDFLMLSNTQFIENR
						VYDEEVEEPVLKAEAEKTEQE
						KTREQKEVDLIPKVQEAVNYG
			1			LQVLDSAFEQLDIKAGNSDSEE
						DDANGRVELILEPKDLYIDRPLP
				1		YLIGSKLFMEQEDVGLGELSSE
						EGSVGSDRGSIVDTE
5718	36086	Α	5764	1	972	

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SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/340,217	sequence	bi peptide sequence	determin, v possone nacreotide insertion)
5719	36087	A	5765	1	2193	MVPYSLLVTWLQKALGVRQY
		1				HVASVLCQRAKVAMRPFEPKY
						IHYDLLEKNINIVCK*LNRPLTL
						SEKIVYRHPDDPTSQEIERGGTY
		1				LQQWPDSVTMQD/ATAQMAM
		1				LQFISSRL/SKVTMPSTIHCDHLI
						EAQLGSEKDLHQAKDINHEVY
						NFLATGSVKYGLSFWKPGSGII
						HQIILENYEYPGVLLIGTDSHTP
						NGGSLGGICIGVGGA/DAVDVM
						TGIPWELKCPKVIGVKLMGSLS
		1				GWTSPKDVILKVAGILTMKGG
}						TGAIVEYHGPVVDSISCTGMVT
						VCNMGLEIGTTTSVFHYNLRM
						KKYMSKTCQADIANLADEFKD
		ĺ				HLVPDPGCHYDQLIEVNPSELK
		İ				LHINGPFTPDLAHPVAEVGKVA
		l		1		EKEEWPLDI*VGLWGSCTNSSY
						EDMGCSAGVAKQALAHGLKC
		1				KSQFTITPCSEQIRASIEQDGYA
						QTLRDVGGIVLANACGPCIGQL
						HRKDINKQEKNIIVTSYNRNFM
						GHNDTNPETHAFITSPETVTAL
						AIVGTLKFNPETNYLTGKDGKK
						FKLEAPDADELPQAEFGPGQDT
						YQHP\PWGTTDHISAAGP*LKFR
						GHLDTISNNLLSGAINIENGKAN
ŀ						SVHDAVTHEFGPFPDTARYKK
						YGISWVVIGD*NYGKGSSWEH
						AALEPRHL\GGRAIITKSFARIHE
						TNLRRQGLLPLTFADPADNKIH
						LVDKLTIQSLKDLT/PGKPLKYI
5720	36088	В	5766	3	1205	
5721	36089	Α	5767	435	1428	
5722	36090	Α	5768	1	615	
5723	36091	Α	5769	79	381	DKPQPHLQHTRTSKRLNRSSQA
						FLQNLLPQELATSAGNLAIRPR
						NACSLGFLVSRVPSLAATPRAP
						WNSGPRLSD*LLPRSSRLSS*RL
		L				TLPDRLGSPVDH
5724	36092	A	5770	38	452	
5725	36093	Α	5771	435	524	
5726	36094	A	5772	284	411	
5727	36095	A	5773	1	354	
5728	36096	В	5774	1	942	
5729	36097	Α	5775	1	417	
5730	36098	Α	5776	88	279	EDLGRSQSESLGPEFQG*VVAIE
		1				TIWPTEPKIFTICLSGSHSQSPW
1		1				NSGPRLSD*LLPRSSQLSS*RLT
				<u> </u>		LPNRLRSLQDHHRCSR

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence		in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5731	36099	Α	5777	206	816	FIDSMLSPSRYQ*LSSQNWKKL
						L*ISYGTKNEPA*PKQS*AQRTK
						LEASH*LTSNCTTSLH*SKQHGT
						GTKTEKKT*AIPFRT*AWAKTS*
į						LKHQKQWQQKPK*TKKIKYLG
	1		ŀ			IQLTTDVKDLFKENYKPLLNEI
						KEDTNKWKNIPCSWIGGINIMK
		l				MAILPKVIYRFNALPIKIPMTFF
ļ						TELEKTTLNFIWNQKRARIAKTI
						LSTKNKAGGITLADFKLYYKPT LVKTAWYWYQNREENLGNTIQ
						DIGMGKNFMTKTPKAMATKA
						KIDKRDLIKPKSFCIAKETTIRM
						NRQPTEWEKLFAIYPSDKALIS
						GIYKDLKQIYKKKNNPIKKWA
						KEMNRHFSKEDIYAANRHVKN
İ						AHHHWSSEKCKSKPQ
5732	36100	A	5778	502	600	
5733	36101	A	5779	51	452	ADKQLQQSLRIQNQCTKITSILI
						HQQQTNREPNHE*TPIHNCFKE
	1					NKIPRNPTYKGCEGPLQGELQT
		l				TAQGNKKGYKQMEEHSMLMG
						RKNQYRENGHTAQELEKTKVH
		İ				MEPKKSPHRQVNPKPKEQSWR
						ннтт
5734	36102	Α	5780	381	528	LPPPLPAGLC**DQQQNSNAPLI
						H*EKERPPDDSYRIFLPRYHFHD
						DRI
5735	36103	Α	5781	1	477	
5736	36104	Α	5782	180	496	TPIHNCYKENKIPGNPTYKGCE
			ļ		•	GPLQGELQTTAQ*NKRGQKQM
						EEHSMLMDRKNKYHEN/EPYSP
	1					R*FIDSMPSPSNYQ*LSSQNWK
5727	26105	ļ	5702	1	522	KLL*GPYGTKKEPT*PRQS MGPEICTSDKLSDEADVTVLEV
5737	36105	Α	5783	1	533	
						LARAIRQEKEIKGIQLGKEEVKL SLFADDMIVYLENPIVSAQNLL
						KLISNFSKVSGYKINVQKSQAF
						LYTNNRQTESQIMSELPFTIASK
						RIKYLGIQLTRDVKDLFKENYK
	-					PLLKEIKEEHSIKWKNIPSSWVG
						RISIMKTAILPKVYIRIKDDEWN
						IYRRYTEFRSLHHKLQNKYPQV
						RAYNFPPKKAIGNKEAVLCHG
						R*HDCISRKPHRLSPKSP*ADKQ
						LQQSLRIQNQCAKITSILIYQ*QT
						NREPNHE*TPIHNCFKENKIPRN
		Ī				PTYKGCEGPLQGELQTTAQGN
						KRGTFHQMEEHSILMGRKNQY
				1		HENGHTAQGLHPDKRR
L	<u> </u>	1	<u> </u>	t	L	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5738	36106	A	5784	599	870	TPANQQHYRRMTLQLKENQTN RKRQQKQHQQKRPHKNPIQRS ATSKIKGDGSIQP*MLPWGQSP QKALTLATAPEQRKDTPPVRGL SLQD
5739	36107	С	5785	1	1170	
5740	36108	A	5786	317	960	KWPYCPRQFIDSMPSPSSYQRL SSQNWKKLL*SSYGTKKEPTSP SQS*AKRTKLEASRYLTSNYTT RLQ*PKQHGTGTKTEI*INGTQQ SPQK*/QPHVYNYLIFNKPEKKK QWGKDSLFNKWDMDEVGNHH SQQTITRTGNQTPHVFTHRWEL NNEITWTQGAGHHTPGPVVGL GAGRGIALREIPNANDELMVQQ TNMAHIYLCIKPARCADVP
5741	36109	A	5787	903	1050	LPPPLPAGLC**DQQQNSNAPLI H*EKERPPDDSYRIFLPRYHFHD DRI
5742	36110	A	5788	281	491	SFGYIPSNGMAGSNGISSSRSLR NRHTDFHNG*TSLQSHQQCKSV PISPHPLQHLLFPDFLMIAILTGV R
5743	36111	Α	5789	1	381	
5744	36112	Α	5790	1	3231	
5745	36113	A	5791	1312	1463	KGSATCSPPQANATTTTALQML GGIPWGNACGIFTRPPPQTYLN THLSTR*RPLLLLLKCVR*KGSA TCSPPQANATTTTALQMLGGIP WGNACGIFTRPPPQTYLNTHLS TR
5746	36114	A	5792	811	1065	IS*VSFMSITH**WKKKEQQLLG CWLA*M*SMLICV*RERI*THKL E*LIILCI*RMKKILEIQKGMFKL LPY*TKRIMLK/TLNRQLNSTVS SLHSRVDSLEKSNTKLIEELAIA KNNIIKLQEENHQLRSENKLIL MKTQQHLEVTKVDVETELQTY KHSRQGLDEMYNEARRQLRDE SQLRQDVENELAVQVSMKHEI ELAMKLLEKDIHEKQDTLIGLR QQLEEVKAINIEMYQKLQGSED GLKEKKEIIVRLYWSALCG
5747	36115	Α	5793	1	1353	
5748	36116	A	5794	78	503	TKELLHSKRNYHQSEQATYRM GENFCNLLI*QRANIQNLQ*TQT NLQEKNKQPHQKVGEGYQQTL LKRRHLCSQKTHEKMLIITGHQ GNTNQNHNEIPSHTS*NGDH*K VRKKQGHG*SWKPSFSANYRK DKKPNTSCSHSQ

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5749	36117	A	5795	2	200	YGLMIRTIPVDPTIRTTSVSLTV RLATMNPRSRPAPVDT\APGPP HLHKNQKSSQHSQYLDLTSYC
5750	36118	Α	5796	76	195	
5751	36119	Α	5797	1	2445	
5752	36120	A	5798	207	475	TNLQEKNKQPHQKVGKGYEQT LLKRKHSCSQQTYEKMLIITGH QRNANQNHNDIPSHSS*NNDH* KDRKQQMLERM*RNRNAFTLL VGV
5753	36121	В	5799	1	681	
5754	36122	Α	5800	1	798	
5755	36123	С	5801	78	245	
5756	36124	Α	5802	16	491	EPYPDRPREPKRDPEWSWGLW DKGGL\VLSLGRTRTEAHTALS RLRASMWIDRSTRAVSVHFTLY NPPTQLFTSVSLRVEILPTGSLV PSSLVESFSIFRSDSALQYHLML PQLVFLALSLIHLCVQLYRMMD KGVLSYWRKPRNWLEVASLVS FSFEK
5757	36125	Α	5803	3	2791	
5758	36126	Α	5804	219	392	
5759	36127	Α	5805	1	1044	
5760	36128	Α	5806	1	360	·
5761	36129	Α	5807	1	528	
5762	36130	Α	5808	127	360	
5763	36131	A	5809		421	GRRSRLAMRPLSMSYSFDLSDV TTPESTKNLVESSMVNGGLTSQ TKENGLSTSQQVPAQRKKLLR APTLAELDSSESEEETLHKSTSS SSVSPSFP\EEPVL\EAVFTRKKP PKFLPISSTPQPERRQPPQRRHSI EKETP
5764	36132	Α	5810	328	1086	
5765	36133	A	5811	1	2076	
5766	36134	Α	5812	1	450	
5767	36135	Α	5813	901	1506	
5768	36136	С	5814	17	106	
5769	36137	A	5815	24	330	ARDQPI/PKKGTVEGFEPADNK CLLRATDGKKKISTVVSSKEVN KFQMAYSNLLRANMDGLKKR DKKNKTKKTKTSSSSSSSSPA AATAATTAATTAATAAQ
5770	36138	Α	5816	1	2247	
5771	36139	A	5817	1	339	
5772	36140	Α	5818	3	463	
5773	36141	Α	5819	3	83	
5774	36142	Α	5820	ī	642	
5775	36143	Α	5821	2	583	
5776	36144	Α	5822	193	409	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5777	36145	A	5823	3	425	
5778	36146	A	5824	1	450	
5779	36147	В	5825	82	2096	
5780	36148	В	5826	1	1554	
5781	36149	A	5827	129	522	LCTDNIEITSKVHLHCSHHGAG
3761	30149	A	19627	129	322	LGTCSPLCLSLPPPPWAPVQPEP PR*APPPAPQRPVPSTIQGRKRL QLHS*SQ*DH/DAHAKVCSFTPE PARPRTHQKEETPNTSEHQKKQ TPDTPPLRTVTLTARLHGF
5782	36150	Α	5828	1	541	
5783	36151	Α	5829	1	714	
5784	36152	A	5830	188	287	KELPSGGHYNCRAPSSPHPAGS S*SGHQPNSQQ
5785	36153	A	5831	71	689	SPVLDDCPQGPLPSEESW/EQPV TRQYPLRPEVHKGLQNIVKHLK APGLVRKCSSHCNTPILGVQKP NGQWRLVQDLRLINEAVVPLY PVVPNPYTLLSQTPEEAEWFMV LDLKDAFFCIPLHSDSQFLFAFE DPTDHTSQITWMVLPQGFRDSP YLFGQALAQDLGHFSSRGTLVL QYADDLLFATSSEASCQQATLD LLNFLANQ
5786	36154	В	5832	45	1148	
5787	36155	С	5833	233	319	
5788	36156	С	5834	11	133	
5789	36157	A	5835	215	849	IKRLPLMKRMWL*LQPESLEIA GILVQ*MIE*QPKKGTNSIPVSK PSP/VVQKPNGQWRQVQDLRLI SDAVIPLYPAVSNPYTLLSQILE EAEWFTVLDLKDAFFCIPLRSD SQFLLAFEDPTDHTSQLTWTVF TQGFMDTPHLFGQSLAQDLGH FSSPGTLVLQYVDDLLLAKQQA TLDLLNFLANQGYKLSKLKAQ LCELLVFSSCARMHS
5790	36158	A	5836		449	MNKELGISSYKFCGMGRKTSVS AAKSVSATIPISRVQGLLQVLG QEVFLLLCQDQEEQAKREKRD QRKATALAMALRQTNLGGSER IENGAGQSPSRACYQCGLQGHF KKDYPTR/EPAAPCPLC*GNHW KMHCPRGQRSCDSHLAITHL
5791	36159	A	5837	1	723	
5792	36160	Α	5838	186	602	
5793	36161	Α	5839	l	843	
5794	36162	В	5840	1	1779	
5795	36163	Α	5841	1	1182	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5796	36164	A	5842	23	302	LSQWKNRWTRTFLASLSVIANI LEPLLNILHVAVWLMLPN*S*V KKQVIM\FWNWWVLGLTDFKN EATDPRVCSWPCWLRSEAADL
5707	26165	_	5042	1	2622	CGECYSS
5797	36165	В	5843	279	2622 1024	
5798 5799	36166	A	5844 5845		3619	
	36167	A		3	1898	
5800	36168	В	5846	1		
5801	36169	A	5847	1	420	
5802 5803	36170 36171	A	5848 5849	1	1632 597	KYLTKKTQAASVEAVKMLDEI
						LLQLSASVPVDVMPGEFDPTNY TLPQQPLHPCMFP/LATAYSTLQ LVTNPYQATIDGVRFLGTSGQN VSDIFRYSSMEDHLEILE*LRVR HISPTAPDTLGCYPFYFCGNTPS FGSKIIRGPEDQTVLLVTVPDFS ATQTACLVNLRS/LACQPISFSG FGAEDDDLGGLGWAPDSKKW
5804	36172	A	5850		1490	RDAARRALNLLGLGEAVWTKQ VRSVAMFSEQAAQRAHTLLSPP SANNATFARVPVATYTNSSQPF RLGERSFSRQYAHIYATRLIQM RPFLENRAQQHWGSGVGVKKL CELQPEEKCCVVGTLFKAMPL QPSILREVSEEHNLLPQPPRSKY IHPDDELVLEDELQRIKLKGTID VSKLVTGTVLAVFGSVRDDGK FLVEDYCFADLAPQKPAPPLDT DRFVLLVSGLGLGGGGGESLLG TQLLVDVVTGQLGDEGEQCSA AHVSRVILAGNLLSHSTQSRDSI NKAKYLTKKTQAASVEAVKM LDEILLQLSASVPVDVMPGEVFD PTNYTLPQQPLHP\CMFPLATA YSTLQLVTNPYQATIDGVRFLG TSGQNVSDIFRYSSMEDHLEILE WTLRVRHISPTAPDTLGCYPFY KTDPFIFPECPHVYFCGNTPSFG SKIIRGPEDQTVLLVTVPDFSAT QTACLVNLRSLACQPISFSGFG AEDDDLGGLGLGP
5805	36173	В	5851	13	441	
5806	36174	Α	5852	3	279	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5807	36175	A	5853	23	841	EKAKEFRRAEEKKKEVPAVPET LKKK*RNFAELKIKRLRKKFA QKMLQRARRKFIHEKAKH\YH KEYRQMYRT\EILMA\RMARKA GNFYVPAEPKLAFVIRIRGINGV SPKVRK\VLQLLRLRQIFNGNLL *SSTKAS\INMLRDCRSHILAWG VPQILKSVNELIYK\RGYGKINK KRIA\LTDN\ALIARSLGK\YGIL LAWEDLIHEIYTV\GK\RFKEAN NFPVGPSNLSSPPRWK*KKKTT HFCKKVEDAGNREGRTRSNRP YLEGMDLRCLP
5808	36176	С	5854	57	287	
5809	36177	Α	5855	1	450	
5810	36178	Α	5856	1	549	
5811	36179	A	5857	3	581	EQRKIPLVPENLLKKRKAYQAL KATQAKQALLAKKEQKKGKG LRFKRLESFLHDSWRQKRDKV RLRRLEVKPHALELPDKHSLAF VVRIERIDGVSLLVQRTIARLRL KKIFSGVFVKVTPQNLKMLRIV EPYVTWGFPNLKSVRELILKRG QA\KVKNKTIPLTDNTVIEEHLG ECYSLGVSWGRKPGSLKL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
}	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
5812	36180	A	5858		2652	MFGRSRSWVGGGHGKTSRNIH
3012		ľ``	5050			SLDHLKYLYHVLTKNTTVTEQ
		E				NRNLLVETIRSITEILIWGDQND
					!	SSVFDFFLEKNMFVFFLNILROK
						SGRYVCVQLLQTLNILFENISHE
						TSLYYLLSNNYVNSIIVHKFDFS
						DEEIMAYYISFLKTLSLKLNNH
						TVHFFYNEHTNDFALYTEAIKF
						FNHPESMVRIAVRTITLNVYKV
						DNQAMLHYIRDKTAVPYFSNL
						VWFIGSHVIELDDCVQTDEEHR
						NRGKLSDLVAEHLDHLHYLND
						ILIINCEFLNDVLTDHLLNRLFLP
						LYVYSLENQDKVFLIIHHAPLV
						NSLAEVILNGDLSEMYAKTEQD
						IQRSSVLPTLSSLWQGSHLSLNQ
						LQSGLHKCSSHLCGAQAAADS
						VTGEIPAIRSLEWLISAGSKART
						FFFLKMLIGFWEKVDCEYQRR
						QVLSTRLQEALPSNRLTDVAAV
						HSSCMLGFGSTAPRGSWIGDPA
						AVHLPLPGELAEHLGSKGTTTV
						TKHQPQAKPSIRCFIKPTETLER
						SLEMNKHKGKRRVQKRPNYK
						NVGEEEDEEKGPTEDAQEDAE
						KAKGTEGGSKGIKTSGESEEIE
	-					MVIMERSKLSELAASTSVQEQN
						TTDEEKSAAATCSESTQWSRPF
						LDMVYHALDSPDDDYHALFVL
						CLLYAMSHNKENLLKKRKAYQ
						ALKATQAKQELLAKKEQKKGK
						GLRFKLLESFLHDYWQQKPDK
5813	36181	Α	5859	1	405	LEISIMAASISGYTFSAVCFHSA
						NSNADHEGFLLGEVRQEETFSIS
						DSQISNTEFLQVI\KVIGWYRFR
						RNTQQQMSYREQVLHKQLTRI
						LGVPDLVFLLFSFISTANNSTHA
						LEYVLFRPNRRYNQRISLAIPN
5814	36182	В	5860	1	843	

			SEQ ID NO:	I .		Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hođ	in USSN 09/540,217	location of first	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		09/340,217	sequence	or peptiae sequence	determine a possible indetermine insertion)
5815	36183	A	5861	3	1260	 EISIMAASISGYTFSAVCFHSAN
3613	30163	^	3001]	1200	SNADHEGFLLGEVRQEETFSISD
						SQISNTEFLQVIQIYNHQPCSKL
						FSFYDYASKVNEESLDRILKDR
		ŀ				RKKVIGWYRFRRNTQQQMSYR
						EQVLHKQLT\RILGVPDLVFLLF
						SFISTANNSTHALEYVLFRPNRR
						YNQRISLAIPNLGNTSQQEYKV
						SSVPNTSQSYAKVIKEHGTDFF
						DKDGVMKDIRAIYQVYNALQE
						KVQAVCADVEKSERVVESCQA
			•			EVNKLRRQITQRKNEKEQERRL
						QQAVLSRQMPSESLDPAFSPRM
						PSSGFAAEGRSTLGDAEASDPP
						PPYSDFHPNNQESTLSHSRMER
						SVFMPRPQAVGSSNYASTSAGL
			ĺ			KYPGSGADLPPPQRAAGDSGE
						DSDDSDYENLIDPTEPSNSEYSH
						SKDSRPMAHPDEDPRNTQTSQI
5816	36184	A	5862	1	742	
5817	36185	A	5863	1	3063	MSLEVDRSVETMCSGDEILLPD
						LPKADVADPLWGPFPVQNCLS
						LARSDSREQGLVLVMESRNRE
						VVPPGVSYSKDGAKSLKGDVP
						ASEVTSKDSTFSQFSPISSAEEC
						GDDEKIKVDDPLTRRTCNQASG
						SAPQQDYDKLKAFGGENSSKT
						GLSPSGNMEKNKVVKREAEAN
				ŀ		SINLSVYEPFKVRKAEDKLKEN
						SDNVLENRVLDGKLSSEKNDT
						CLPGTAPSKTKSSSKLSSCSSAI
						MALSAKKAASDSCKEPV
5818	36186	Α	5864	2	626	LCQHNRIKHKGIRKVYACSHCP
						DSRRTFTKRLMLEKHVQLMHS
			:			LKDPDLKE/TDRCHQ*GGNRNK
		ļ				RRH/LSSPVPSGSWKNQFSR/CA
1						MCGFTTENLLQFHEHIPQHKSD
1		ļ				SSSYQCQECGLWYTSHVSLSRH
]]					LFIVHKLKEPQPVSKQNGAGED
						NQQENKPSHEDESPNGTVSDRK
}						CKVCVKTFETEAALNTHMRIH
		<u> </u>	l	<u> </u>	<u> </u>	GMAFIKSKRMSSAKK

NO:	of peptide sequence	hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide	codon for last amino acid	*=Stop codon, /=possible nucleotide
5819	36187	L		sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
		Α	5865	1	765	MQRDVYISHVRKEHGKQMKK
						QPCCQRDKPFSSSHSLCWHNRI
						KHKVIRKVYTCSHCSDSRGTFT
1 1	i					KQLMLEKHVQLTHGIKDPDLK
						E/TDRRHQ*GGNRNKRRPQGPQ
						SQVEVERTGSGVQASQ\GAITQ
	;					PLKKLKINVFKVHKCARA\GFT
						TENLLQFHEHIPQHKSDGSSYQ
						CREHLFVVHKLKEPQPVSKQN
						GAGEDNQQENKPSHEDESPNG
						AMSDRKCKVCAKTFETEAALN
						THMQTHGMAFIKSKRMSSDEK
5820	36188	Α	5866	878	1095	AFFLDTLYTQFSRC*SRAFVPFA
						HPTAMACSSRR/LIRVQEQDAA
						QVTHQSFKDDLCVVLGDTLHH
						RFFLIFRH
5821	36189	Α	5867	1	1137	
5822	36190	Α	5868	1	1380	
5823	36191	Α	5869	55	558	
5824	36192	Α	5870	1	897	
5825	36193	A	5871	-1	565	
5826	36194	В	5872	22	1696	
5827	36195	Α	5873	295	522	
5828	36196	A	5874	3	3678	IPAPVSGRPPPGLLAEGKLSGPR
						PMCRTRLGSHTAASAPARMWL
						FHTLLCIASLALLAAFNVDVAR
ļ						PWLTPKGGAPFVLSSLLHQDPS
						TNQTWLLVTSPRTKRTPGPLHR
						CSLVQDEILCHPVEHVPIPKGRH
			•			RGVTVVRSHHGVLICIQVLVRR
						PHSLSSELTGTCSLLGPDLRPQA
						QANFFDLENLLDPDARVDTGD
			İ			CYSNKEGGGEDDVNTARQRRA
]]						LEKEEEDKEEEDEEEEAGT
						EIAIILDGSGSID
5829	36197	A	5875	1	3189	2

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
5830	36198	Α	5876	1119	2463	LRAGGPGQGGKGPAHPGRRVD
						ELLLGHPTAVGCAKGTDCPTPT
			•			PGGLWV*GVQEESSDLPTAVDS
						SRPDIRDQAWASVHWELYVHG
						SSFINT*GERGAGY/AVITW/STV
						VEARSMPQGTSAQKAELIAFIR
						ALELSEALAKTVRQRCVSCRQ
						HHARQGPAVPPGIQAYGAAPFE
		l				DLQVDFTEMPKCGDIRKIVTGD
						VNTPAILGVVSSSPPSHIGNNITE
						DPELQPILAGLSLSMYLVTVLR
						NLLIILAVSSDPHLHTPMCFFLS
						NLCWADIGFTLATVPKMIVDM
						QSHTRVISYEGCLTRISFLVLFA
						CIEDMLLTVMAYDCFVAICRPL
						HYPVIVNPHLCVFFLLVYFFLSL
						LDSQLHSWIVLQFTIIKNVEISN
						FVCDPSQLLKLACSDSVINSIFM
						YFHSTMFGFLPISGILLSYYKIVP
		l				SILRISSSDGKYKAFSTCGSHLA
5831	36199	В	5877	1	1830	
5832	36200	С	5878	109	245	
5833	36201	В	5879	1	801	
5834	36202	В	5880	171	288	
5835	36203	В	5881	49	502	
5836	36204	В	5882	1	639	
5837	36205	Α	5883	1	821	MSGQFDFLLWTPEEDNSDESD
	,					AEGEHGDGAEEEAPPPVPRPGP
]				KPAGLGRRPCPYEQAQGGDGP
						EEQWMSFCWDIPRKSGAPRERI
						TLRRHLEDCGCKVSKKKAQICR
1						QQKDPQEESPAQFYERLCEAYG
						MYTPFDPDSPENQRMIHMALV
		1				RQSAEDMRRKLQKQAGLAGM
						NPSQLLERASQVFVNRDAV/TP*
						GKQQREWRERG*YDSQYRRGG
						TQPCDIVPNIRRERR*YHSQYRR
						RARPPGPH*SERGALRPGSLPRQ
						AA*RVPSGPADGQ
5838	36206	Α	5884		1860	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
5839	36207	A	5885	2745	3732	IRIGKNYFKVHMEPKKSPHRQV NPKPKEQSWRHHTT*LQTILQG YSNQNSMPGPTPQ\PSTPAPGG NLRNPQSSDLLQVTKQQGQAL AIQREAPLHRIPAPEAIPWYFQP QPATQLGSPPVDPPSSAMM/SR RAHRSRGPDRQGYPLQGD*PGE PRPQEH*RGHSQERLPSSEKQTP ICPPAQATQHPEEPDAHQPYKH
						LFQVCAHQGHPVAQRRN*PGY WERYHSAEDPELQPILAGLSLS MYLVTVLRNLLIS
5840	36208	В	5886	1	2379	
5841	36209	В	5887	1	6741	
5842	36210	В	5888	165	928	
5843	36211	Α	5889	56	2028	
5844	36212	В	5890	1335	3804	
5845	36213	Α	5891	365	1573	
5846	36214	Α	5892	1	1491	
5847	36215	Α	5893	25	458	
5848	36216	A	5894		1194	MGGNAADKFRAGVELALQSG NKVCVCSPRTTISGDAVAPRSV LPGCRHIPGEISQHSHCWAGRS LSSRKRGSLFRRMGFIKVVKNK AYFKRYQVKFRRR\RKGKT\DY YARKR\LVIQDKNKYNTPKY\R MIV\RVTNRDI\ICQMLYA/RRIE GDMIVCAAYCTPNLPKIWV*RV GLTNYAAAY\CTGLL\ARRLL NRFGMDKIYEGQV\ELTGDEYN VESIDGQPGAFTCYLDAGLAR\ TTTGNKVFGAPEGKLWMGGLS IPHSTK\RFPG\YDSE\SKEFNA\E VTSGKHIMGPKCLQNYM\RYL ME\ED\EDAYKK\QFS\QYIK\NS VT\PDMM\EEMYK\KAHAAIRE ESSSMEKKAQGKKFKKKRWNR P\KMSLAQKKDRVAQKKASFL QSSRSGLLESLTPAIFP
5849	36217	Α	5895	1	1191	
5850	36218	A	5896	21	159	PRSSVRPAGGFSSTCPPSWAAS RSCTTSHGCGHQ*CAGTGTQTR L
5851	36219	A	5897	1	1062	
5852	36220	Α	5898	1	246	
5853	36221	Α	5899	1	642	
5854	36222	A	5900	2	648	
5855	36223	Α	5901	1	555	
5856	36224	В	5902	151	513	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5857	36225	A	5903		2915	VSLAFCQPLSLSLSPLLPLASSL APERTHLPGPGSLLLSPPSFPAR PREPRGCVTAAPPDKMDTAEE DICRVCRSEGTPEKPLYHPCVC TGSIKFIHQECLVQWLKHSRKE YCELCKHRFAFTPIYSPDMPSRL PIQDIFAGLVTSIGTAIRYWFHY TLVAFARWEVVLLPACRIYKCL FLGSVSSLLTLPLDMLSRKNLL ADCLQGCF\VVTCTLCAFISLV WLREQIVHGGAPIWLEHAAPPF NAAGHHQNEA
5858	36226	Α	5904	3	907	
5859	36227	В	5905	102	406	
5860	36228	В	5906	1	5241	
5861	36229	A	5907	1	519	
5862	36230	A	5908	1	3067	TRDAAMAEAALEAVRSELREF PAAARELCVPLAVPYLDKPPTP LHFYRDWVCPNRPCIIRNALQH WPALQKWSLPYFRATVGSTEV SVAVTPDGYADAVRGDRFMM PAERRLPLSFVLDVLEGRAQHP GVLYVQKQCSNLPSELPQLLPD LESHVPWASEALGKMPDAVNF WLGEAAAVTSLHKDHYENLYC VVSGEKHFLFHPPSDRPFIPYEL YTPATYQLTEEGTFKVVDEEA MEKAEVSRTCLLTVRVLQA
5863	36231	Α	5909	1	429	
5864	36232	Α	5910	1	690	
5865	36233	Α	5911	1	376	
5866	36234	A	5912	1	471	
5867 5868	36235	A	5913 5914	3	2756	SPGGRTPAARDSIVREVIQNSKE VLSLLQEKNPAFKPVLAIIQAG DDNLMQEINQNLAEEAGLNITH ICLPPDSSEAEIIDEILKINEDTRV HGLALQISENLFSNKVLNALKP EKDVDGVTDINLGKLVRGDAH ECFVSPVAKAVIELLEKSGVNL DGKKILVVGAHGSLEAALQCLF QRKGSMTMSIQWKTRQLQSKL HEADIVVLGSPKPEEIPLTWIQP GTTVLNCSHDFLSGKVGCGSPR IHFGGLIEED
5869	36237	A	5915	3	404	RLKEKKLVKEVIAVSCGPAQCQ ETIRTALAMGADRGIHVEVPPA EAERLGPLQVARV/AVVTADLR LNEPCYATLPNIMKAKKKKIEV IKPGDLGVDLTSKLSVISVEDPP QRTAGVKVETTEDLVAKLKEIG

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5870	36238	Α	5916	1	334	
5871	36239	A	5917	88	870	MADVRVLVAVKRVIDYAVKIR VKPDRTGVVTDGVKHSMNPFC EIAVEEAVRLKEKKLVKEVIAV\ SCGPAQCQETI\RTALA\MGA\D RGIHVE\VPPAEAERF/GVPLQV ARVLGQAWQRRRRLDLVLLGK QA\IACTDCNQ\TGQMTAGFLD WPQGTF\ASQVTLEGDKLKVER EIDGGLETLRLKLPAVVTADLR LNEPRYA\TLP\NIMKAKKKIE VIKPGDLGVDLTSKLSVISVEDP PQRTAGVKVETTEDLVAKLKEI
5872	36240	Α	5918	1	774	
5873	36241	Α	5919	1	1557	
5874	36242	Α	5920	1	759	
5875	36243	Α	5921	305	638	
5876	36244	Α	5922	364	540	
5877	36245	Α	5923	9	111	
5878	36246	Α	5924	1	594	
5879	36247	Α	5925	169	594	
5880	36248	Α	5926	37	936	
5881	36249	В	5927	91	2043	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	l .	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
5882	36250	Α	5928	1	2298	MSRKGPRAEVCADCSAPDPGW
						ASISRGVLVCDECCSVHRSLGR
		s.				HISIVKHLRHSAWPPTLLQMVH
						TLASNGANSIWEHSLLDPAQVQ
			ļ			SGRRKANPQDKVHPIKSEFIRA
						KYQMLAFVHKLSCRD\DDGVT
						AKDL\SKAN*HSSVR\TGKPWK
						TCLRLLS\LGA\QANFFHPEKGT
						TPLHVAAKAGQTLQAEL\LVVY
						GADPGSPDVNGRTPIDYA\RQA
						GHHELAERLVEC\QYE\LTERLA
						FYLCGRKPDHKNGHYIIPQMAD
						SLDLSELAKAAKKKLQALSNRL
						FEELAMDVYDEVDRRENDAV
						WLATQNHSTLVTERSAVPFLPV
						NPEYSATRNQGRQKLARFNAR
		j				EFATLIIDILSEAKRRQQGKSLSS
						PTDNLELSLRSQSDLDDQHDYD
						SVASDEDTDQEPLRSTGATRSN
						RARSMDSSDLSDGAVTLQEYL
						ELKKALATSEAKVQQLMKVNS
	:					SLSDELRRLQREIHKLQAENLQ
]			LRQPPGPVPTPPLPSERAEHTPM
		l				APGGSTHRRDRQAFSMYEPGS
ŀ						ALKPFGGPPGDELTTRLQPFHS
		1				TELEDDAIYSVHVPAGLYRIRK
						GVSASAVPFTPSSPLLSCSQEGS
						RHTSKLSRHGSGADSDYENTQS
		l				GDPLLGLEGKRFLELGKEEDFH
						PELESLDGDLDPGLPSTEDVILK
						TEQVTKNIQELLRAAQEFKHDS
		l				FVPCSEKIHLAVTEMASLFPKRP
5883	36251	A	5929	1	924	I VI CSERIILA VI EMASLI I RIG
5884	36252	A	5930	214	387	RRAFHPTPVTSARSTLHSFGQL
3004	30232	^	3930	214	367	RLPHSM*SGGTLLISMMQANQS
						LSLEFSKWTSFPLD
5885	36253	Α	5931	1	1639	LOLEFORWISFFLD
5886	36254	A	5932	1	1098	
5887	36255	A	5933	1	546	NSSQVLQF*YLWVNEAPFRLLT
3007	30233	ľ		•		MPQPYPQINITADQLWGSVAVL
						V*MH\NGLPGLCI*AWEKTTSG
						GEQYPSSSAIKQGPKEPYADFIA
						RLQESLKKVIADLAAQDIVLWL
						LAFDNANPECQAALRPIRGKAH
						LVDYTKVCDGIRDKLHKATLL
						AQAMAGLRMGKGNTPFPGACF
				-		NCGKHGH
	1	<u> </u>			<u></u>	PACOKHOTI

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
5888	36256	IA	5934	<u> </u>	648	MGQVWGLVHFTLEVFHTGDEE
						EQEYSEVTEDVTEHVYLPAKA
		ŀ				KVAKEEEAGIQQARQEGDLEA
	1					WQFPVRIHPPDQQENITATFEPF
						PFKLLKELKQAINQYGPGSPFV
						MGLLKNVTVSSQMIPTDGDPLT
		İ				RACLTPAQFLQFKTWWADEAS
						IQAARNAWAQPQINITADQLLG
						VGGWAGLDAQFVMQDDAIEQ
						LRGVCIRAREK\IT*CGEQYPSF
5889	36257	Α	5935	120	254	
5890	36258	C	5936	366	479	
5891	36259	Α	5937	1	933	
5892	36260	A	5938	1	840	MGKVWGLVHSTLELFHTDVEE
						EGEYNEVTEEVTKHVYLPAKA
						KAAKEGEVHPYHSAPPHYYFE
						ENDPPDLSFLEDTGTPRLAAAA
						TTKANA\KQPLLLSPSMHR/PLP
E						TRPLLAG\QATALPQGV*PLIAP
						PTRPHLEHCSTQ*RPQPAPTLGE
						ITPVTVPQAPLHPRAVQPW/HST
						PTRSWLQAVMPGIAPPSRSPPW
						AVQLRILTYLGCLSTTPAAVQS
						LSQPPASVRQARVPQAPASSLC
						QEKPPSPPPVAPHGAAAARPP
	<u> </u>					SYHSSCASVAVVAANVLIG
5893	36261	Α	5939	1	134	MGQVW*KSQSPKA*EPGVSCP
						RAGRREASVPRWYSS
5894	36262	Α	5940	1	1158	
5895	36263	Α	5941	1	673	MGIALGSPTTGPKAMKQNLSIG
						PAGILLPQPVMTPGCPAAAPPV
						APRCGYVLSEDAWGNLQGLCT
						GLSCGQPVAPWATDSGGAQDP
		1				VALLPPFPSRGLLGPSCMQRSQ
į.						VVKGGRQSEISSQNKKVIADLA
						AQDIVLQLLAFDNANPDCQAA
			1			LRPIRGKAHLVDFIKACDVIGGI
						PNLLLHPRGLRAITIAVFGKQ\IR
						TSGSNPLRSTFLSRSRSISKNCSA
5005	26264	-	5040	ļ.	1100	EV
5896	36264	Α	5942] [1182	L

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
5897	36265	A	5943	1056	2128	TQPPTWGQIKKLSQMVEENLR
						KAGQLVTMNNLMVAMIVVITT
						TEEQEYSEVTEEVTEQVYLPAK
						AKVAKEGEVHPYPSAPPHYYFE
			ŀ			EKEWPDPPDLSFLEDTROKVVA
						PVTVRAAPRATVLSSIQAGIQQ
						AR*EGDLEAWQFPVRIHPPDQQ
		ŀ				GNIIVTFEPFPFKLFKEFKQAVN
		ļ				QYGPGSPFVMGLLKNVAVSSW
						MIPTDWDALTRACLTPAQFLQF
						KTCAMKQGPREPYVDFIARLQE
						SLKKMIADSAAQEIVLQLLAFD
						NAHPDCQAALRPIRGKAHLVD
			ļ			
						YVKACDGIGEYTPSTYEKKVN
						CKTASCRSFRRYPEEGIVITGDD
						SSMRVVVPEDLPVGQDVEVED
5000	26266	<u> </u>	5014	ļ 1.	200	RDIDDPDPV
5898	36266	Α	5944	11	300	
5899	36267	Α	5945	1	373	
5900	36268	Α	5946	1	2976	
5901	36269	A	5947	287	544	
5902	36270	A	5948	831	4555	
5903	36271	C	5949	108	434	
5904	36272	C	5950	145	402	CORPORISE COMPONIA COMP
5905	36273	A	5951	3	1104	GSRPGGHTLQSHRGRHAQGLR
		l				SPLLAPVCLGCGTWGQRRLFW
	1					SLKI*RLLCWF/CRRLQINCKNK
		ŀ				PAIPRGPT/GPSEKSGLLLQDQG
						DTTNTEEENSKSLENIFGRVIKE
			ŀ			NFQSLVRDLDI\QYKKHKEHPG
						NSSQKDHHLGTLSSGYPKLRQR
		l				KES*ELSDRSTRKISQS/YNLYTP
1		ŀ	ł			NTGAPRFIKQLLTDLRNEIDSNT
		ŀ				IIVGYFSTPLTALDRSSRQKVND
İ		ŀ				ETVDLNYTLEQMDLTEIYRTFH
		l				PTTTEYTFYSTVHGTFSKIDHMI
						CHKMSLNEFKKTEIISSTLSDHS
	ĺ		1			GIKLEINSKMNLQNHANTWKL
		1			1	NNLLLNEHWVKNEIKMEIEKFF
					[KLNDSNDTTYQNLWDMAKAV
						LRGKFIALNTYIKKMM
5906	36274	В	5952	1	2367	
5907	36275	Α	5953	21	132	WREGARQCQEDPAEFLSD*DW
		<u> </u>				ADPGPPHNAPGQNWH
5908	36276	Α	5954	l	1461	

sequence 09/540,217 codon for peptide of peptide sequence deletion, \=pti 5909 36277 A 5955 31 802 SAWWNS *EYPPTSC LRMNGDO	n, /=possible nucleotide ossible nucleotide insertion) RELLDPQSRSRSGFCF
*EYPPTSO LRMNGD	RELLDPOSRSRSGEGE
*EYPPTSO LRMNGD	
LRMNGD	GTEQESVSGNRALCSS
1 1 1 1 1	QNSDVYAQEKQDFV
	RVLTEDEMGHPEIGD
1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	VLEYNAIGGKYNRGL
1 1 1 1 1 1 1	RELVEPRKQDADSLO
1 1 1 1 1 1	WCVELLQAFFLVAD
1 1 1 1 1	TRRGQICWYQKPGV
	DANLLEACIYRLLKLY
1 1 1 1 1	/LNLIELFLQSSYQIEI
	PHSPPRAMWILSDSLK
	SSTRQLSTPSTFL
	OKYKDIQDKLQIISEET
1 1 1 1 1	CMALKADVVAKKRA
1	LYNRSLNEYKALKKD
	RIEELKK/RVLTN/RLE
1 1 1 1 1 1 1 1 1	KKISWLKERVKAFQN
1	QEIEQFQQALEKDKEE
1 1 1 1 1 1	EKLDVKHALSYNQRL
1 1 1 1 1 1	SKTDRLKRFGPNVPAL
LEAIDD	
5911 36279 A 5957 121 475	
5912 36280 A 5958 131 2891	
5913 36281 A 5959 131 3101	
5914 36282 A 5960 44 292 CQDYRW	LLASSVLCWLY*KTQ
QSDTEDL	LCSAPTGPPNPEEDD
GKS*PEK	CRQMT*KKWSINCKC
FFASSHN	TTLSIGLFLR
5915 36283 B 5961 26 239	
5916 36284 A 5962 131 3121 LVSFLLFG	QDALPEGDASPLGPY
LLPSGAP	ERGSPGKEHPEERVV
TAPPSSSC	QSAEVLGELVLDGTA
PSAHHDI	PALSPLLPEEARPKHA
LPPKKKL	PSLKQVNSARKQLRP
KATSAAT	ΓVQRAGSQPASQGLD
LLSSSTER	KPGPPGDPDPIVASEE
ASEVPLW	VLDRKESAVPTTPAPL
QISPFTSQ)PYVAHTLPQRPEPGE
PGPDMAC	QEAPQEDTSPMALMD
KGENELT	rgsaseesqetttstiit
TTVITTEC	QAP
5917 36285 A 5963 1 1185	
50.0	NVSFNHWSAIFLTME\
5918 36286 A 5964 495 636 PSVVLDI	CT 1 1 FOR 4 CO
	SHAMESR*GSVLP*N
	SHAMESR*GSVLP*N

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence		*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
		<u> </u>	ļ	<u> </u>		
5920	36288	Α	5966	3	688	GTSAGDGCPHPHTVLWPPDIMP
						PKKDVPVKKPAGPSISKPAAKP
		-				AAAGAPPAKTKAEPAVPQAPQ
						KTQEPPVDLSKVVIEFNKDQLE
						EFKEAFELFDRVGDGKILYSQC
						GDVMRALGQNPTNAEVLKVLG
						NPKSDEMNVKVLDFEHFLPML
						QTVAKNKDQGTYEDYVEGLRV FDKEGNGTVMGAEIRHVLVTL
						GEKMTEEEVEMLVAGHEDSNG
						CINY\EELVRMVLNG
5921	36289	A	5967	54	540	AVYMCDLTENQTADLIPST\EF
3721	30207	` `	3701			KEA\FQLFDRTGDGKILYSQCG
						DVMRALGQNPTNAEVLKVLGN
						PKSDEMNVKVLDFEHFLP\MLQ
						TVAKNKDQGTYEDYVEGLRVF
		ŀ				DKEGNGTVMGAEIRHVLVTLG\
		1				EKMTE\EDVHMLVAGHEDSNG
						CINYEAFVRHILSG
5922	36290	A	5968	1	405	
5923	36291	Α	5969	93	506	
5924	36292	В	5970	43	615	
5925	36293	Α	5971	2	598	TRTSLHAALFSWDPVGTSLETFI
						SHLEVKKGMLVGIVGKVGCGK
						SSLLAAIAGELHRWEQVQCGG
		-				HTLH*APEQMLRGHVAVRGLS
			1			KGFGLATQEPWIQFATIRDNILF
						GKTFDAQLYKEVLEACALNDD
						LSILPAGDQTEVGEKGVTLSGG
						QRARIALARAVYQEKELYLLD
5926	36294	В	5972	137	371	DPLAAVDADVANPPACTGCNP
5927	36295	A	5973	1137	1398	
5928	36296	A	5974	1	915	
5929	36297	A	5975	40	564	
5930	36298	A	5976	62	280	
5931	36299	A	5977	17	744	
5932	36300	Α	5978	1	537	
5933	36301	Α	5979	242	576	AANSLYKGSNSAALIILVGP*IS
						CLTHPVYMK*APS*TICLLYSM
		l				VILKALFFPQNINFIFLPVKSRHS
5934	36302	Α	5980	924	3473	
5935	36303	Α	5981	1	604	
5936	36304	Α	5982	288	732	RNCIANYDFLILPTSF*WCSFFV
				}		LLFLFLSLSLSSLKLHSDLDKGE
						GTAKYTLSGDGAGP\VFTIDETT
						GD\IHAIRSL\DREEKPFYTLR\A
						Q\AVDIETRKPLEPESEF\IIKVQ
					!	DI\NDNEPKFWGWDLMVATGP
	2 62 0 7				1.5.0	QKWSPVG*VGQIKIL
5937	36305	Α	5983	1	1510	L

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	1	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5938	36306	Α	5984	1	603	
5939	36307	A	5985	250	505	GFSPTWQ*MNTHPSLMCFSPSR TTGSAAAN*RGQQKSGVPFWS* SHVKGKDQIPVPICPCTTAPVT MESTCIC*PTHVSCKTLMV
5940	36308	A	5986	3	131	RVLVPDIRGAI*AVPKSPSIKSYP ALPRALFLLPPYPDDLGI
5941	36309	Α	5987	1	913	
5942	36310	A	5988	5	177	
5943	36311	Α	5989	3	383	
5944	36312	A	5990	1	1323	
5945	36313	Α	5991	342	594	
5946	36314	Α	5992	1	321	
5947	36315	Α	5993	3	601	
5948	36316	A	5994	1	1023	
5949	36317	Α	5995	1	606	
5950	36318	Α	5996	1	1536	
5951	36319	Α	5997	2	1394	
5952	36320	Α	5998	371	1045	
5953	36321	Α	5999	3	360	
5954	36322	Α	6000	1	636	
5955	36323	С	6001	1	861	
5956	36324	Α	6002	1	711	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
5957	36325	A	6003	118	2446	AGSPPRADAAAGLWERGHLLS
3,5,	30323	` `				AAVAAMGKRDRADRDKKKSR
						KRHYEDEEDDEEDAPGNDPOE
		ł				AVPSAAGKQVDESGTKVDEYG
						AKDYRLQMPLKDDHTSRPLWV
						APDGHIFLEAFSPVYKYAODFL
		ŀ				VAIAEPVCRPTHVHEYKLTAYS
		ĺ				LYAAVSVGLQTSDITEYLRKLS
						KTGVPDGIMQFIKLCTVSYGKV
						KLVLKHNRYFVESCHPDVIQHL
						LQDPVIRECRLRNSEGEATELIT
						ETFTSKSAISKTAESSGGPSTSR
						VTDPQGKSDIPMDLFDFYEQM
		İ		·		DKDEEEEEETQTVSFEVKQEMI
						EELQKRCIHLEYPLLAEYDFRN
						DSVNP\DINIDLKPTAV\LRPYQE
		1				KSLR\KMFGNGRARSGVIVLPC
						GAGKSLVGVTAACTVRKRCLV
						LGNSAVSVEQWKAQFKMWSTI
						DDSQICRFTSDAKDKPIGCSVAI
						STYSMLGHTTKRSWEAERVME
						WLKTQEWGLMILDEVHTIPAK
						MFRRVLTIVQAHCKLGLTATLV
						REDDKIVDLNFLIGPKLYEANW
						MELQNNGYIAKVQCAEVWCP
						MSPEFYREYVAIKTKKRILLYT
		•				MNPNKFRACQFLIKFHERRNDK
						IIVFADNVFALKEYAIRLNKPYI
						YGPTSQGERMQILQNFKHNPKI
		I				NTIFISKVGDTSFDLPEANVLIQI
]						SSHGGSRRQEAQRLGRVLRAK
		1				KGMVAEEYNAFFYSLVSQDTQ
5958	36326	A	6004	1	933	
5959	36327	A	6005	1	879	
5960	36328	A	6006	1	727	
5961	36329	A	6007	1	1509	
5962	36330	A	6008	1	1698	
3702	20220	<u> </u>	10000	1	11070	<u> </u>

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
5963	36331	A	6009	2465	2751	VPTSSAPMPWPSRQWERSSRT MTVISSSQLMALGP*RTAQVLA AAPA*LGASSAARGIDPQAGKT QK\ETDIATMQLCANKLDKKDF FGKSDPFLVFYRSNEDGTFTICH KTEVVKNTLNPVWQPFSIPVRA LCNGDYDRTVKIDVYDWDRD GSHDFIGEFTTSYRELSKAQNQ FTVYEVLNPRKKCKKKKYVNS GTVTLLSFSVDSEFTFVDYIKGG TQLNFTVAIDFTASNGETRMSE KVGGNPLQPTSLHYMSPYQLS AYAMALKAVGEIIQDYDSDKL FPAYGFGAKLPPEGRISHQFPLN NNDEDPNCAGIEGVLESYFQSL RTVQLYGPTYFAPVINQVASLP
5964	36332	Α	6010	1	1059	
5965	36333	Α	6011	255	405	
5966	36334	Α	6012	251	1199	
5967	36335	Α	6013	1	1791	
5968	36336	A	6014	335	984	
5969	36337	A	6015	1	127	LPLRLSYEIYLYFTF*KPEHNKS HFLESHLIRNTRSALNTRK
5970	36338	Α	6016	1	774	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
5971	36339	A	6017	186	2699	PRGAKPAVPAGPERPGPGPC
İ						SPRPMVLPTCPMAEFALPRHSA
		ł				VMERLRRRIELCRRHHSTCEAR
						YEAVSPERLELERQHTFALHQR
					ļ	CIQAKAKRAGKHRQPPAATAP
						APAAPAPRLDAADGPEHGRPA
						THLHDTVKRNLDSATSPQNGD
						QQNGYGDLFPGHKKTRREAPL
		ŀ				GVAISSNGLPPASPLGQSDKPSG
						ADALQSSGKHSLGLDSLNKKR
!	1					LADSSLHLNGGSNPSESFPLSLN
						KELKQEPVEDLPCMITGTVGSIS
						QSNLMPDLNLNEQEWKELIEEL
						NRSVPDEDMKDLFNEDFEEKK
						DPESSGSATQTPLAQDINIKTEF
		l				SPAAFEQEQLGSPQVRAGSAGQ
						TFLGPSSAPVSTDSPSLGGSQTL
						FHTSGQPRADNPSPNLMPASAQ
1		1				AQNAQRALAGVVLPSQGPGGA
						SELSSAHQLQQIAAKQKREQML
						QNPQQATPAPAPGQMSTWQQT
						GPSHSSLDVPYPMEKPASPSSY
						KQDFTNSKL/PHDA*CE*EFPSA
		İ				RRPLPPAQPCEPAESPATE*LES
		ł				ELRE/YTRGLCW\IRQYKTPFSL
						QSGLWARQPGVWPEQASPDGL
						SSPAAVPYKSRAELPVSDEAKA
						RKYAFPITGSTWPGAEPFQCPC
		İ				ASPGYQCWDPAACRVRGQLPQ
						QLPLSQQPATGRCNEAASVAFG
						PTETKGAAAKAFTATAVPSEAT
		İ				APSRGTGEATVSAPSDPPTTPVP
5972	36340	Α	6018	1	1062	
5973	36341	Α	6019	3	566	ASRPPVTLRLVVPASQCGSLIG
						KGGCKIKEIRESTGAQVQVAGD
:						MLPNSTERAITIAGIPQSIIECVK
•						QICVVMLESPPKGVTIPYRPKPS
						SSPVIFAGGQAYTIQGQYAIPQP
						DLTKLHQL\AMQQC\HFPMTHG
						\NTGFQWALES\SFPEV\KGYW\
		1				AGLDASAQTTSHELTIPNDLIGC
		L				IIGRQGAKIN
5974	36342	Α	6020	2	531	TLRSSWQVVRTGTAQAATVRG
						FPPHPPRP\SASTLTLEGPPLEAY
						TIQGQYAIPQPDLTKLHQL\AM
		}				QQSHFPMH/GDGIHSDYSAG\LD
]			ASA\QTTSH\ELTISKRILIGLA*S
						GRSRAPKINGGSRQMSGAQI\KI
						ANPV\EGSTD*GRFTITGSAASIS
L		1	<u></u>			LAQYLINVRLSSETGGMGSS

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	I .	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
5975	36343	Α	6021	181	1369	PLKVQLPRTLLDMDTGVIEGGL
						NVTLTIRLLMHGKEVGSII\GKK
					•	GESVKKMREESGARI\NISEGNC
:	ŀ					PERII\TLAGTH*LPSFKAFAMI\I
						DKLGRRDISSSM\TNSTAAQLGP
						PVTLKLG\APAS\QCGSLIGKGG
						CKIKEIRESTGAQVQVAGDMLP
						N\STEAGHHYLLGIPQSII\ECV\K
						QILRGSCWETLFPSSPPKGVTIP\
						YRPK\PSSSPV\IF\AGGPGQVPA
						QAATVRAFPTPPRPCGSTLTWE
						GPPLEAYTIQGQYAIPQPD\LTK
						L\HQL\AMQQSHFPMTHGNTGF
						SG\IESSF\PEVKGYW\AGL\DAS
	ŀ					l i
	ļ					AQTTSHELTI\PNGFDWAGIIGA
						FKGAKIN\EIRQMSGAQIKIANP
						VE\GSTDRQVTITGSAASISLAQ
5076	26244		(022	1	(00	YLINVRLSSETG\GMGNN
5976 5977	36344	A	6022 6023	1	690 399	
	36345	A		1	397	
5978 5979	36346 36347	A	6024 6025	3	249	
		A A	6026	177	477	
5980 5981	36348 36349	A	6027	49	451	
5982	36350	A	6028	ļ	1305	
5983		-	6029	1	568	
	36351	A	6030	207	1053	PLTFGPARWRETPPSLYKEFSG
5984	36352	Α	10030	207	1033	l l
						LFGSFFLSSAWGAHNLRALFLL
	1				1	LPSSRIQPTPWVFGDLKSPAGL
	1					QVLNDYLADKSYIEGYVPSQA\
						DVAVFEAVSSPPPADLCHALR
						WYNHIKSYEK\EKASLPGVKKA
						LGKYGPADVEDTTGSGATDSK
						DDDDIDLFGSDDEE\ESEKPKRL
						\REERLAQYESKKAKKPG\LVA
						KSSILLLVKP\WD\DETDMAKLE
						ECVRSIQADGLVWGSSKL\VPV
						GYGIKKLQIQCVVEDDKVGTD
						MLEEQITAFEDYVQSMDVAAF
5985	36353	A	6031	1	211	
5986	36354	Α	6032	27	241	EVFGSWWQVPHEWLGSVLVV
						MSEFLFH*LM*ELVLKRV*CLS
	1					HLSLAPFLAM*YACSSSSPSTMI
		<u> </u>				VSFFFF
5987	36355	Α	6033	1	588	

Sequence	op codon, /=possible nucleotide on, \=possible nucleotide insertion)
	, , , , , , , , , , , , , , , , , , , ,
GTE GDL SVT	
GTE GDL SVT	CGCWLFASLTVMEAAHFFE
GDL SVT	KLLEVWFSRQQPDANQGS
SVT	RTIPRSEWDILLKDVQCSII
	•
	KTDKQEAYVLSESSMFVSK
1 1 1 1 1	ILKTCGTTLLLKALVPLLKL
	OYSGFDSIQSFFYSRKNFMK
1 1 1 1 1 1	QGYPHRNFQEEIEFLNAIFP
	AGYCMGRMNSDCWYLYTL
	EER\VISQPDQTLEILMSELD
	MDQF*HEDG\VTAKDVTRE
	RDLIPGSVIDATMF\NPCGYS
	GMKS\DGTYWT\IHITPEPEF
	SFETNLSQTSYDDLIRKVV
1 1 1 1 1 1 1	KPGKFVTTLFVNQSSKCRT
	LPQK\IEGFKR\LDCQEGWF
	NFFF\TSFAKKQQQQQS
5989 36357 A 6035 I 687	-
5990 36358 A 6036 1 894	
5991 36359 B 6037 I 1458	
5992 36360 A 6038 I 1062	
5993 36361 A 6039 I 1218	
	RIAWAQEMDAAGRGCHLL
1 1 1 1 1	AARGPARAPAAAAAAAAS
i i i i i i i i i i	PCSGAACAPSAAAGAGAM
	SSAGEEKGATGGSSSSGSG
1 1 1 1 1 1	CCLGAEGGADPRGAGSAA
	GAAALDEPAAAGQKEKDEA
	KLRNLTFRKQVSYRKAISR
	.QHLAPAHPLSLPVANGPA
	RATLDWSENAVNGEHLWL
	VSG\DLCYLGEENCQVRFA
	LRRKCAVCKIVV/HTTACIE
	KINFRCKPTFREG\GS
5995 36363 A 6041 45 155	
5996 36364 A 6042 I 979	
5997 36365 A 6043 3 1042	
5998 36366 A 6044 2 401	
5999 36367 A 6045 1 5772	CDL I EVEDEN CDCCCL I I D
1	GRLLFYFRSMGPGCDLLLR
1 1 1 1 1	TITAAAIMSDTDSDEDSAG
	PFSLAGFLFGNINGAGQLEG
	LDDECKKHLAGLGALGLG
	TELTANEELTGTDGALVND
	VVRSTEDAVDYSDINEVAE
	RRYQQTMGSLQPLCHSDY
	DYDADCEDIDCKLMPPPPP
	PMKKDKDQDSITGEKVDFS
	DSESEMGPQEATQAESEDG
1 1 1 1 1	LPLAGIMQHDATKLLPSVT
ELFI	PEFRPGKVLRFLR
6001 36369 A 6047 1 354	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
6002	36370	Α	6048	1	666	
6003	36371	Α	6049	129	456	TYPAVGESSRRPERSRGVPGPD
						AAAESHFRPGRISNPLHRAGEQ
	1					GHARAEGSGVFRPHRGRGLRL
	1					LFVQAPDQVDAPVHG*VAPPAP
						GARDAQTCRSHECPRTRPLDEV
6004	36372	Α	6050	1	1095	
6005	36373	A	6051	171	619	
6006	36374	A	6052	1	621	
6007	36375	В	6053	1	181	
6008	36376	Α	6054	1	905	
6009	36377	Α	6055	1	855	
6010	36378	Α	6056	2	244	
6011	36379	Α	6057	3	829	LTDCLKDVDLIPPFNRMLLEVT
						FGKLYAWTVQNILNVLMDASA
						KFKELGIQPVPLQTITSENPLGP
						SLGSIPQARFLLMMLSMLTLQH
		l				SANNLDLLLNSGTLALTQTALR
						LIVLPDVELLI*DHLL*MQTSIV*
						LPVLALLLAQHRSFGPSCDNVE
						EDMNASAQGASATVLEATRKE
						TAPVHLPVSGPELAATMKIGTR
•						VMRGVDWKWGD/QEAEQTER
						NIHPTTMMFTSTINLLQTLCLPA
						RVHAEIMQSEATKTLCGLLQIL
						VESGTTDKTRME
6012	36380	Α	6058	133	5421	IFVFGCPGSPRLQMVENRYTES
						FFTGQNCRNNEKVTLVRIADLE
		ŀ	ŀ			NHNNDGGFWTVIDGKVYDIKD
						FQTQSLTENSILAQFAGEDPVV
						ALEAALQFEDTRESMHAFCVG
						QYLEPDQEGVTIPDLGSLSSPLI
			-			DTERNLGLLLGLHASYLAMSTP
			1			LSPVEIECAKWVQSSIFSGGLQT
					Ì	SQIHYRYNEEKDEDHCSSPGGT
						PASKSRLCSHRRALGDHSQAFL
						QAIADNNIQDHNVKDFLCQIER
			1			YCROCHLTTPIMF

SEQ ID NO:	of peptide	Met hod	SEQ ID NO: in USSN	location of first	codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nuclcotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
6013	36381	Α	6059	1	2607	MMGHSSAIPLTATPGELKGQSP
						TKMPDPELGCQGAK\$QGCSRN
						ARHQKARSMPLQDQHLALAIL
						LELAVQRGTLSQMLSAILLLLQ
						LWDSRAQETDNERSAQGTSTL
						LLSLLQTFQSIICSKDTPPSEGN
						MHLLSGPLSPSESFLRESFFTVQ
						NCRNNEEVTLICKADLENHNK
l E						DGGFWIVIDEKVYDIKDFQTQS
		į				LTGNSILAQFAGENPVVALEAA
				: :		FEFEVTRESMHAFCVGQYLEVR
						LYALSDAEDGRG\TL*WLQSSIF
			<u> </u>			SG/GLQTSQIHYSYNEEKDEDH
						CS/SPVGTPASKSR\CSHRWALG
						DHSQAFLQAIADNNIQDHNVKT
						HQEQGRSYKEVCTPVIERLRFL
						SNELRPAVGNDLSIISEFKLLSSL
						PRWRRIAQKIIRERRKKRIPKKP
	,					ESTADEEKIGNEESDLEEACILP
				1		HSPINVDKRPIAIKSPKTITSENP
						LGPSLGSIPQARFLLMMLSMLT
						LQHSANNLDLLLNSGTLALTQT
						ALRLIGPSCDNVEEDMNASAQ
						GVSATVLEATRKETAPVHLPVS
						GPELAATMKIGTRVMRGVDW
						KWGDQDGPPPGLGRVIGELGE
				i		DGWIRVQWGTGSTNSYRMGK
						EGKYDLKLAELPAAAQPSAEDS
						DTEDDSASPNRLVYREQHRSW
						CMLGFVRSIALTPQVCGALSSP
]						QWITLLMKVMKGHAPFPAASL
						QRQRWVAVSLPHALVKSGTVP
6014	26202	_	(0(0		7007	QRQRWVAVSLITIALVRSGTVF
6014 6015	36382	A	6060	32	7297 259	SLRVTPGGRNVSEKKDRKKTG
0013	36383	A	10001	32	239	i i
						R/PPELERKSHTERLYLEEGSAE
}						PA*VGRRAPESVNDRGGAIQRA
6016	26204	 	(0(2	2	127	QRGQPAPPLPR
6016	36384	A	6062	3	437	MTELDTI WIZDI IZMEDATEDALAD
6017	36385	Α	6063	¹	425	MTELPTLWKPLKNSPMFRVIAR
						AFCVTFILVSTHQEMLTITSENP
						LGPSLGSIPQARFLLMMLSMLT
						LQHSANNLDLLLNSGTLALTQT
						ALRLIGGTPVELPAELAAPAFR
						AAAA/VDKNPRRRGQKVAVAK
L		L	<u> </u>			SGGDRGKKSR

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
6018	36386	A	6064	2	14566	RRGGRALRVRPETWEEAGEKM PSESFCLAAQARLDSKWLKTDI QLAFTRDGLCGLWNEMVKDG EIVYTGTESTQNGELPPRKDDS VEPSGTKKEDLNDKEKKDEEET PAPIYRAKSILDSWVWGKQPDV NELKECLSVLVKEQQALAVQS ATTTLSALRLKQRLVILERYFIA LNRTVFQENVKVKWKSSGISLP PVDKKSSRPAGKGVEGLARVG SRAALSFAFAFLRRAWRSGEDA DLCSELLQESLDALRAL
6019	36387	В	6065	75	581	
6020	36388	A	6066	517	5495	ESFFTGQNCRNNEKVTFVRIAD LENHNNDGGFWTVIDGKVYDI KDFQTQSLTENSILAQFAGEDP VVALEAALQFEDTRESMHAFC VGQYLEPDQEGVTIPDLGSLSSP LIDTERNLGLLLGLHASYLAMS TPLSPVEIECAKWLQSSIFSGGL QTSQIHYSYYEEKDEDHCSSPG GTPASKSRLCSHRRALGDHSQA FLQAIADNNIQDHNVKDFLCQI ERYCRQCHLTTPIMFPPEHPVEE VGRLLLCCLLKH
6021	36389	Α	6067	1	1851	
6022	36390	A	6068	53	423	YHLTLLIHGTKVSATRPGHCSP GKLPSWVPISLQCTPSSPECRSV *RQL*HGHMHELLYIITKADSVI TASCLISRSEW\WGDGGTVSLV QRFVDYGSSQCQRPHRTGCASS SAPQRASSVAPG
6023	36391	Α	6069	1	49	EWSGTRTWWRPWSCRT*CRT
6024	36392	A	6070	1	726	
6025	36393	A	6071	368	485	
6026	36394 36395	A	6073 6074	3	535	RRREFPAKERESASMSSPEE*P/DMAKGAPPSQPPPNSVPAD/CSRSVRLSSSNFEESLRPTEDDEGQEGQDSMSRAKANWLRAFNKVRMQLQEARGEGEMSKSLWFKGGPGGGLIIIDSMPDIRKRKPIPLVSDLAMSLVQSRKAAITSALASSTLNNEELKNHVYKKTLQALIYPISC
6028	36396	Α	6075	1	222	
6029	36397	A	6076	2	284	NGIPDFSNTTLLRGLQMPGARG PCCGSQRHINLLMGTGVFLVK MKA/NEFFFLDSETRLCKIAPEG WREQPQKTSMNTFTLFLRIKFF VSHYGLL

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
6030	36398	Α	6077	91	749	
6031	36399	С	6078	1	380	
6032	36400	A	6079		877	MPRPDLAPALLVAVLVLLYLL VSAAAPAPSSDSTPENVCPVET TTNFIWELSSPHNTSAVLLVVF HKGPSPKFEVKLKKNANGLGFS FVQMEKESCSHLKSDLVRIKRL FPGHPAEENGAIAAGDIILAVN GSPRKASSSRCRGSWAMQLSV QAGPSFASYYPAAVEVLHLLRG APQEVTLLLCRPPPGALPELEQE WQGQKYKCAWAIVNEAEVEE VVEVAGTVEFFGDQ\GA*CHHP ACSGKPLGWHGGLSRKCPRLFF RGATRVTGAQGLKEVKQQSFR SNMMWLGLRL
(022	26401	_	6000	1	1602	SINIVINIWEGERE
6033	36401	A	6080	1	1683	
6034	36402	В	6081	66	1214	
6035 6036	36403 36404	A	6082	3	5694 483	SSLTSSMEDPAAPGTGGPPANG
						NGN\GGGKGKQAAPKGREAFR SQRRESEGSVDCPTLEFEYGDA DGHAAELSELYSYTENLEFTNN RRCFEEDFKTQVQGKEWLELE EDAQKAYIMGLLDRLEVVSRE RRLKAARAVLYLAQGTFGECD SEVDVLHWSRY
6037	36405	Α	6084	1	2646	
6038	36406	A	6085	115	1289	FSPLEPRLCSLGGWGALQAGEP CQPSRAGCGREGATMGCTLSA EERAALERSKAIEKNLKEDGIS AAKDVKLLLLGAGESGKSTIVK QMKIIH\EDGFSGEDVKQYKPV V\YSNTIQSL\AAIVRAMDTLGIE YGDKERKADAKMVCDVVSRM EDTEPFSAELLSAMMRLWGDS GIQECFNRSREYQLNDSAKYYL DSLDRIGAADYQPTEQDILRTR VKTTGIVETHFTFKNLHFRLFD VGGQRSERKKWIHCFEDVTAII FCVALSGYDQVLHEDETTNRM HESLMLFDSICNNKFFIDTSIILF LNKKDLFGEKIKKSPLTICFPEY TGPNTYEDAAAYIQAQFESKNR SPNKEIYCHMTCATDTNNIQVV FDAVTDIIIANNLRGCGLY
6039	36407	Α	6086	1	1053	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
6040	36408	Α	6087	2	377	IYDGFDGFAKGQIKEIGWTDVG
				_		GWTGQGGSILGTKRVLPGK/Y
		Ì				MEEIATQMRTHSINALLIIGGFE
						AYLGLLELSVFVTLKLFYQHFM
		1	•			HVLLTLIPNRRVPICPTSPSACC
		1				LWSVSHAFRCAYEQN
6041	36409	Α	6088	1	591	
6042	36410	Α	6089	1	2391	PGRAAPLGLLAMDADDSRAPK
		}				GSLRKFLEHLSGAGKAIGVLTS
		1				GGDAQGMNAAVRAVVRMGIY
						VGAKVYFIYEGYQGMVDGGSN
						IAEADWESVSSILQVGGTIIGSA
			}			RCQAFRTREGRLKAACNLLQR
						GITNLCVIGGDGSLTGANLFRK
						EWSGLLEELARNGQIDKEAVQ
						KYAYLNVVGMVGSIDNDFCGT
	1					DMTIGTDSALHRIIEVVDAIMTT
Ì		•			į	AQSHQRTFVLEVMGRHCGYLA
						LVSALACGADWVFLPESPPEEG
		l				WEEQMCVKLSENRARKKRLNII
						IVAEGAIDTQNKPITSEKIKELV
						VTQLGYDTRVTILGHVQRGGTP
l		ł	ł			SAFDRILASRMGVEAVIALLEA
			<u> </u>			TPDTPACVVSLNGNHAVRLPL
	1	ł				MECVQMTQDVQKAMDERRFQ
						DAVRLRGRSFAGNLNTYKRLAI
						KLPDDQIPKTNCNVAVINVGAP
						AAGMNAAVRSAVRVGIADGH
1						RMLAIYDGFDGFAKGQIKEIGW
						TDVGGWTGQGGSILGTKRVLP
]	GKYLEEIATQMRTHSI\NALLIIG
						GF\EAYLGLLELSAAREKHEEFC
<u> </u>	l'					VPMVMVPATVSNNVPGSDFSIG
		ŀ				ADTALNTITDTCDRIKQSASGT
1						KRRVFIIETMGGYCGYLANMG
						GLAAGADAAYIFEEPFDIRDLQ
						SNVEHLTEKMKTTIQRGLVLRN
						ESCSENYTTDFIYQLYSEEGKG
[VFDCRKNVLGHMQQGGAPSPF
6043	36411	A	6090	1	459	
0043	120411	Ι^_	10090	L1	T/	<u> </u>

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
6044	36412	A	6091	1930	5781	RAKSPANIIMTGSNSHITILTLN
						VNGLNSPIKRHRLASWIKSODP
						SVCCIQETHLMCRDTHRLKIKG
						WRKIYQANGKQKKAGVAILVS
						DKTDFKPTKIKRDKEGHYMMV
						KGSIQQEELTILNMYAPNTGAP
						RFIKQVLSDLQRDLDSHTLIMG
						DFNTPLSTLDRSTRQKVNKDTQ
				1		ELNSALHQADLIDIYRTLHPKST
		}				EYTFFSAPHHSYSKIDHILGSEA
						LLSKCKRTEIITNYLSDHSAIKL
						ELRIKNLTQSR
6045	36413	Α	6092	1	3654	
6046	36414	Α	6093	1	5127	
6047	36415	Α	6094	1	3663	
6048	36416	Ā	6095	1	3210	MVKGSIQQEELTILNIYAPNTG
						ALRFIKQVLRDLQRDLDSHTIIM
						GDFHTPLSTLDRSTRQKVNKDI
						QELNSALHQEDLIDIYRTLHPKS
						TEYTFFSAPHHTYSKIDHIVGSK
						ALLSKCKRTEIITNCLSDHSAIK
						LELRIKNLTQNRSTTWKLNNLL
						LNDYWVHNEMKAEIKMFFETN
						ENKDTTYQNLWDTFKAVCRGK
						FIALNAHKRKQERSKIDTLTSQL
						KELEKQEQTHSKASRRQEITKIR
						AELKEIETQ
6049	36417	Α	6096	1	5073	
6050	36418	Α	6097	1	3489	MGDFNTPLSTLDRSTRQKVNK
						DTQELNSALHQADLTDIYRTLH
						PKSTEYTFFSAPHHTYSKIDHIV
						GSKALLSKWKRTEIITNYLSDH
}		İ		ļ		SAIKLELRIKNPTQSRSTTWKLN
ł						NLLLNDYWVHNKMKAEIKMFF
						ETNENKDTTYQNLWDAFKAVC
						RGKFIPLNAHKRKQERSKIDTL
						TSQLKELEKQEQTHSKASRRQE
						ITKIRAELKEIETQKTLQKINESR
						SWFFERINKTDRPLARLIKKKR
						EKNQIDTIKNDK
6051	36419	Α	6098	1	3235	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	1	*=Stop codon, /=possible nucleotide
	sequence	İ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
6052	36420	A	6099	1	3070	MATLIPLLFHKHPHLTQRYQFN
						VRGGCPTSTSAIATLDPRTAVFP
						LMSVFQTSAPWPFWGFTHQTP
						VPQPLKVKIHSSADTAADIQAN
						RVWSGPPANSKRPAAEEGKLT
						NRKDIHTKNPSVRHHHQRPKV
		ŀ				DKTTKTGKKQRRKTGNSKKQS
		ļ		[ACPPPKERSSSTATEQSWTEND
						FDKLREGFRRSNYSKLQDEIQT
						KGKEVENLEKSLDECITRITNTE
						KGLKKLENYVKNEETSGANAI
						NWKKGINKIDRPLARL
6053	36421	Α	6100	1	3297	
6054	36422	Α	6101	1	2563	MKAEIKMFFETNENKDTTNQN
						LWDAFKAEEVESLNRPITGAEI
						GAIINSLPTKKSPGPDGFTAEFY
	Ì	l				QRYKEELVPFLLKLFQSIEKEEI
	į					LPNSFYEASIILIPKPGRDTTKKE
						NFRPISLMNIDAKILNKILANRI
						QQHIKKLIHHDQVGFFPGMQG
;						WFNIRKSINVIQHINRAKDKNH
						MIISIDAEKAFDKIQQPFMLKTL
						NKLGIDGTYFKIIRAIYDKPTAN
						IILNGQKLEAFPLKTGTRQGCPL
	1					SPLLFNILLEVLARAIRQEKEIK
		l				GIQLGKEEVKLSLFADDMIVYL
						ENPIVSAQNLLKLISNFSKVSGY
:						KINVQKSQAFLYTSNRQTESQI
		1				MSELPFTIASKRIKYLGIQLTRD
						VKDLFKENYKPLLKEIKEDTNK
		ł				WKNIPCSWVGRINIVKMAILPK
						VIYRFNAIPIKLPMTFFTELEKTT
						LKFIWNQKRAHITKSILSQKNK
						AGGITLPDFKLYYKATVTKTA
		l				WYCYQNRDIDQWNRTEPSEITP
		İ	i			HTYNYLIFDKPEKNKQWGKDS
						LFNKWCWENWLAIWRKLKLD
						PFLTPYTKINSRWIKDLNVRPKT
						IKTLEENLGITIQDIGMGKDFMS
						RTPKAMATKAKIDKWDLIKLK
1				1		SFCTAKETTIRVNRQPTTWEKIF
						ATYSSDKGLISRIYNELKQIYKK
			[KTNNPIKKWEKDMNRHFSKED
						IYAAKKHMKKCSSSLAIREMQI
			1			KTTMRYHLTPVRMAIIKKSGNN
6055	36423	Α	6102	1	3417	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
6056	36424	Α	6103	1	3579	MPGHNLKWKLNRGTVLIETGI
						QLSTSTILGSASEPPSAPIPKAQV
						SSTEKLRNCIDDLKPFPALASEL
						SRRAKALQIAGFPPMKVPRDTI
					ł	SKVCLDKTVGKLCHSGEESRK
						CTLICNNKHYPIQDNLQGYKTQ
						NKFLNKEILELSALRRNAERRE
	İ					RDLMAKYSSLEAKLCQIESKYL
						ILLQEMKTPVCSEDQGPTREVI
						AQLLEDALQVESQEQPEQAFV
						KPHLVSEYDIYGFRTVPEDDEE
						EKLVAKVRALDLK
6057	36425	Α	6104	1	4371	
6058	36426	Α	6105	1	1825	MVKGSIQQEELTILNTYAAHTG
						APRLIKQVLSDLQRDLDSHTIIM
						GDFNTPLSTLDRSTRQKVNKDT
		-				QELKSALHQADLTDIYRTLHHK
						STEYTFFSAPHHIYSKIDHILGSK
						ALLSKCKRTEIITNYLSDHSAIK
				ľ		LELWIKNLTQNHSTTWELNNLL
						LNDYWVHNEMKAEIKMFFETN
						ENKDTTYHNLWDTFKAVCRGK
				ľ		FIPLNAHKRKQERSKIDTLTSQL
						KELEKQEQTHSKASRRQEITKIR
						AELKEIETQKTLQKINESRSWFF
						ERINKIDRLLARLIKKKREKNQI
						DAIKNDKGDITTDPTEIQTTIRE
						YCKHLYANKLENLEEMDKFLD
						TYTLPRLNQEEVESLNRPITGAE
						IVAIINSLPTKKSPGPDGFTAKF
						YQRYKEELVPFLLKLFQSIEKE
						GILPNSFYEASIILIPKPGRDTTK
						KENFRPISLMNIDAKILNKKLA
İ		1				KRIQQHIKKLIHHDQVGFIPGM
						QGWFNIRKSINVIQHINRAKDK
						NHMIISIDAEKAFDKIQQPFMLK
						TLNKL\GIKYLGIHLTRDVKDLF
						KENYKPLLKEIKEDRNKWKNIP
						CSWVGRINIVKMAILPKNILITL
		<u> </u>				QLLLVLPELSTLIPLWLPALAGQ
6059	36427	A	6106	1	4449	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide inscrtion)
6060	36428	A	6107	824	3693	AWKGTTDRSTRQKVNKDTQEL NSALHQADLIDIYRTLHPKSTE YT/FFSAPHHTYSKIDHIVGSKA LLSKCKRTEIITNYLSDHSAIKL ELRIKNFTQSRSTTWKLNNLLL NDYWVHNEMNAEIKMFFETNE NKDTTYQNLWDAFKAVCRGK FIALNAHKRKQERSKIDTLTSQL KELEKQEQTHSKASRRQEITKIR AELKEIETQKTLQKINESRSWFF ERITKSDRPLARLIKKKREKNQI DTIKNDKGDIT
6061	36429	Α	6108	1	3297	
6062	36430	В	6109	112	3300	
6063	36431	A	6110	3	3316	
6064	36432	Α	6111	1	3457	
6065	36433	Λ	6112		3170	MVKGSIQQEELTILNIYAPNTG APRFIKQVLSDLQRDLDSHTLI MGDFNTPLSTLDRSTRQKVNK DTQELNSALHQADLIDIYRTLH PKSTEYTFFSAPHHTYSKIDHIL GSKALLSKCKRIEIITNYLSDHS AIKLELRIKNLTQSRSTTWKLN NLLNDYWVHNEMKTEIKMFF ETNENKDTTYQNLWDAFKAVC RGKFIALNAYKRKEERSKIDTL TSQLKELEKQEQRHSKPSRRQE ITKMRAELKEIETQ
6066	36434	В	6113	1	4753	
6067	36435	В	6114	1	3384	
6068	36436	Α	6115	1	3345	
6069	36437	Α	6116	1	3780	
6070	36438	Α	6117	1	3720	
6071	36440 36440	A	6118	1	3335	MVKGSIQQEELTILNIYAPNTG APRFIKQVLSDLQRDLDSHTLI MGDFNTPLSTLDRSTRQKVNK DTQELNSALHQADLIDIYRTLH PKSTEYTFFSAPHHTYSKIDHIV GSKALLSKCKRTEIITNYLSDHS AIKLELRIKNLTQSRSTTWKLN NLLNDYWVHNEMKAEIKMFF ETNENKDTTYQNLWDAFKAVC RGKFIALNVYKRKQERSKIDTL TSQLKELEKQEQTHSKASRRQE ITKIRAELKEIETQ
6073	36441	Α	6120	1	3780	
6074	36442	Α	6121	1	3852	
6075	36443	Α	6122	1	3345	
6076	36444	В	6123	1	6136	
6077	36445	В	6124	1	3924	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence	:	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
6078	36446	В	6125	295	4185	
6079	36447	Α	6126	1	405	
6080	36448	Α	6127	1	1059	
6081	36449	A	6128	2	518	EKATCDENDSVSNIATEIKEGQ QSV\HVSPQKQSAWKVIFKKKV SLLNIATRITGGGKSGTVSSQKQ PPSKATSDKTDSALNIATEIKDG LQCGTVSSQKQPALKATTDEED SVSNIATEIKDGEKSGTVSSQKQ PALKATTDEKDSVSNIATEIKD
						GEKSVTVSSQKQPASK
6082	36450	A	6129	1	843	
6083	36451	Α	6130	1	432	
6084	36452	A	6131	279	835	CGSAQAAAAAAAEEATEKIPAL RP/ASAVGAAGALAVLRDPRA WREAGSKS/PETPFS*CQGA/HG GGQFCPSGTAFPGKMKVMMR KRKKKGQCLPGICRSLKRRKSP RSPGM/IGYSTLSIPPEMLASYQ SYSSTFHSLEEQQVCMA/G*HR QMPGADKPVRACQRVLAGVW AGFTMRHQSRLY*ECSICGVIFR HRCVAHLLGRAIGAGLFVITPA GERGMCCKAIKLGNARVFPVT TL*NDGQCQACMHAG/TTLEDP GKNYISSSPSSSSSSSSSPSSPRVPS
6085	36453	C	6133	278	446	
6086	36454	Α	6134	1	1152	
6087	36455	Α	6135	1	639	
6088	36456	Α	6136	12	1436	
6089	36457	Α	6137	367	1056	
6090	36458	Α	6139	l l	4263	
6091	36459	Α	6140	1	1311	
6092	36460	A	6141	1093	1358	MQKKPLTKFNNPSC*KLSIN*V LMGRISK*VTTTATTTTTTTTT TTTTTTTTTTTTTTTTTTTTTTTT
6093	36461	Α	6142	1	699	
6094	36462	Α	6143	1	489	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
6095	36463	A	6144	5	916	ALGGTPMLGKLAMLLWVQQA
						LLALLLPTLLAQGEARRSRNTT
						RPALLRLSDYLLTNYRKGVRPV
						RDWRKPTTVSIDVIVYAILNVD
	1	ļ.				EKNQVLTTYIWYRQYWTDEFL
						QWNPEDFDNITKLSIPTDSIWVP
	İ	1				DILINEFVDVGKSPNIPYVYIRH
						QGEVQNYKPLQVVTACSLDIY
						NFPFDVQNCSLTFTSWLHT/SPG
		l				HQHLFVALARKGEIRQECLHEP
						GRVGVAGGAALLSGVQHGKQ*
		ł				LLCRNEVLCGHPPAAPLLCGQP
						ATAQHLPHGHGHRGLLPAPQQ
		ŀ				WREGLFQNYTPPGLLGLPDHRF
6096	36464	Α	6145	5	756	
6097	36465	Α	6146	1	471	
6098	36466	Α	6147	184	1097	ALGGTPMLGKLAMLLWVQQA
:						LLALLLPTLLAQGEARRSRNTT
			<u> </u>			RPALLRLSDYLLTNYRKGVRPV
1						RDWRKPTTVSIDVIVYAILNVD
						EKNQVLTTYIWYRQYWTDEFL
:						QWNPEDFDNITKLSIPTDSIWVP
						DILINEFVDVGKSPNIPYVYIRH
						QGEVQNYKPLQVVTACSLDIY
						NFPFDVQNCSLTFTSWLHT/SPG
						HQHLFVALARKGEIRQECLHEP
		İ				GRVGVAGGAALLSGVQHGKQ*
		İ				LLCRNEVLCGHPPAAPLLCGQP
						ATAQHLPHGHGHRGLLPAPQQ
						WREGLFQDYTPPGLLGLPDHRF
6099	36467	Α	6148	3	672	
6100	36468	Α	6149	3	398	LRKIKIDLGKFSDNPDGYIDVLQ
						GLRQSFDLTWRDIMLLLDQSLT
						PNEKSAAKTAAREFGDLWFV/S
						FQN*SCKTTNRSSNGAPDAVHD
						*D/PTADPWTGPLAHALMLMTL
						KAPLTRKSQLHDPYYAPIQQEA
L						V
6101	36469	Α	6150	1	420	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence	5	
6102	36470	Α	6151	Ī	878	TLLPPRPGVGSDYPNIPGCPMW
						GNKDPGEGTLERGKVEDASQR
		ŀ				TSRTLKETSGKAEHRKKODGK
					}	CSPQGKNALKMYSGEFDPVRV
						CVPFSLSDLKQLKIDLGRFSDNP
						DGYIDVLQGLGQSFDLTWRDI
						MLLLDQTLTPNERNATITAARE
						FGDLWCLSQVNDRMTTEERER
						FPTGQQAVPSADPH*DTESEHG
						DWCCRHLLTCVLEGLRKTRKK
1						AVNFSVMSTVTQGKEENPTAFL
						ERLREALRKHTSLSPDSIEGQLI
						LKIKFITRSAADIRKQTSKVHLR
						PGAKLKTPY
6103	36471	A	6152	1	417	
6104	36472	Α	6153	1	415	
6105	36473	Α	6154	1	234	
6106	36474	Α	6155	1	335	TCPIVPG/QEMIIEISKGRSGLGL
						SIV*GKDTPLNAIVIHEVYEEGA
			}	,		AAVDGRLWAGDQILEVNGVDL
		-	1			RNSSHEEAITALMQTPQKVRLV
						VYRDEAHYRDEENLEIFPVDLQ
6107	36475	Α	6156	109	4687	
6108	36476	Α	6157	109	4991	
6109	36477	A	6158	3	862	
6110	36478	A	6159	1	1857	
6111	36479	A	6160	3	537	
6112	36480	A	6161	125		
6113	36481	A	6162	125	1184	LSIYLVTQKAA/SSEWGPEQEK
6114	36482	Α	6163	875	1300	ALQEVQAAVQAALILEPYDPA
						GPVVLEVSLADRDAVWSLWQ
						APIGESQQRPLGFWSKALPSSA/
		İ				DHKACHAQQHSIIKWKWYIHD
						RARAGPEGTNSSARYAATM*K
						W\TASALQPLSRKSLKDSSEGK
		1				SSQWAELRAVHLAVHVAWKE
		ŀ				KWPDVRLDTDSWAV/ANGLAR
						WSGTWKEHDRKIGDKEVWGR
						GTRI
6115	36483	A	6164	1	1566	
6116	36484	Α	6165	I	1098	
6117	36485	Α	6166	1	538	
6118	36486	Α	6167	410	625	LTLVPFSFSVVLTPLPKGNTIRY
						*TGIL*GWNLSWKGTLGFSVTA
		1	[QPSSGTPRTALGSRTTKPLLVSL
		1				LQV

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ŀ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
l i				sequence		
6119	36487	A	6168	821	1428	QVTDVQDFPLKCKIQWRRTVR
						GDLQ/PFTRVTVHWGKGNDQT
		1				FQGLLDTGSELTLIPGDPKRHC
						GPPVKIGAYGGQIINGVLAQVQ
						LTVDAVGPWTHPVV/I/SPVPECI
	:					IGIDILSSWQ/NSHIGSLTGRVRA
						TMVGKAKWKPLELPLPRKIVN
						QKQYHIPGGIAEIGATIKDLKDT
						G\VIPTSPFNSPI*PMQKTDGSW
						RMTVDYL
6120	36488	A	6169	203	850	DVEHVGKMRRRWRRRVVVAD
0120	30100	` `	0107			GNVVKPMSCAGDLQ/PFTRVTV
						HWGKGNDQTFQDLLDTGSELT
						LIPGDPKRHCGPPVKIGAYGGQI
						INGVLAQVQLTVDAVGPWTHP
						VV/I/SPVPECIIGIDILSSWQ/NSH
						IGSLTVHLSSDPKGCHSEWGPE
}						QEKALQEVQAAVQAALILEPY
				1		DPAGPVVLEVSLADRDAVWSL
						WQAPIGESQQRPLGFWSKALPS
6121	36489	A	6170	1	864	WQ/II IOESQQIA EGI WSIELEIS
6122	36490	A	6171	1	205	VPDFPAKPIQGPHFNSRYL/REA
0122	30470	l'`	1771		203	VHGEAVHAPFIAGQPLT**ELM
						SIFPQFQFFLYRRLDSHSGQSWQ
						I
6123	36491	Α	6172	1	3039	
6124	36492	Α	6173	1	1364	MIISIDAEKAFDKIQQPFMLKTL
						SKLEETASPSPVVATYTPQPML
						PSAFPPLSEEINPVLPETTVMAS
						PEAVTRQDNVDSPQKPPPTPMF
						ASRPITRLKPRRAPSEEGIQRLK
						KIGMVEWISHFRPTPLSMEGPE
						HILLTNTLLNRYVKAAPASLKS
						PLTALLFMSDLTVGTTFSQLQN
						LNTMGIFGSSCDRSQVAALNHQ
						RQVPELIIGIDILSSWQNPHIGSL
						NGRGYINSLALCHNLIRRDLDR
						FLLPQDITLVHYIDHIMRLDSVK
	1					DKWLHLAPPTTKKEAQCLVGL/
	1					FGFWRQHIS\LWRQPLACYWAL
						VETEHLTMGHQVTMQPELPIM
	1					NWLLSDPSSYKPAPMASWGVP
						YGQLTEEEKTRAWFTDGSARY
					:	TETTRKWTAVAIQPLSRTSLKD
						SNEGKSSHQQAKNGITVLAGVI
						DPDYQDEISL/LTPQWRCHPGSS
						VWRASSLGISHPP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	1			sequence		
6126	36494	ĪA	6175	1	475	MGRNQNRKAENSKNWNASSPP
		` `				KEHNSSPAREQNWREDEFNELT
						EVGFRRSVITNFSDLKEHVLTH
	1					CKEAKILEKRLDKWLTKITSVE
					}	KS/CK*PDG/D*KPQYENFMKHT
						QASIADLIKRKKRYQ*LKIKLM
						K*SKKTR*EKKKYKLPSENTTN
						TSMQIY
6127	36495	A	6176	179	393	NKRGYKQMEEHSMLMDKKNQ
						YYENGHTAQGN*AGEGNKGYS
1	:				1	IRKTGSQIVPLCR*HDCISRKPD
						CLSPKSP
6128	36496	A	6177	1	710	MARELRDECTSFSSQFHQLEER
						VSVIEDQMNEMKQEEKFREKGI
						KRSEQSLQEIWDYMERPNLCLI
						GVPESDGENGIKLEDTLQDIIQK
	1					DFPNLGRQANIQIQEIQRMPQR
						YSSRRATPRHIIVRFTKEIQTTIR
						EYYKHLYANKLENLEEMDKFL
						DTYTLPRLNQEEVESLNRPITGS
						EIEAIINSLPTKKSPRPDGFTAEF
						YQRYKEELVPFLLKLFQSIEKE
	•					GILPNSFYEASIILIPKLGRDTTK
						KENFRPISLMNIDAKILNKILAN
						QMQQNIKNLIRHDQAGFIPGMQ
						CWFNIRKSINVIQHINRTKDKN
						HMIISIDAEKAFYKIQQPFILKTL
						NKLENKIPRNPTYKGCEGPLFK
						ENYKLMLKEIKEDTSKWKNIP
						WSWIGRINIVKMAILPKVIYRLS
						AIPIKLPMIFFTELEKTTLKFIWN
						QKRACIAKSILSQKNKAGGIML
						PDFKLYYKATVTKAAWYWYQ
						NRDIDQWNRTEPSETIPHIYSHL
		1				IFDKPDKNKKWGKDFLFNKWC
						WENWLAICRKLKLDPFLTPYTK
						INSRWIKDLNVRPKTIKTLEENL
						GNTIQDIGMGKDFMCKTPKAM
						VTKAKIDKWDLIKLKSFCTAKE
						TTVRVNRQPTEWEKIFAIYSSD
1					1	KRLISRIYKELKL*PCSIV*SQVA
		\bot				*CLQLCSFGLGLTWQCRLFFGSI
6129	36497	Α	6178	464	724	IGEVLLDNILQRVFQLGSILPITF
		<u> </u>				RYTNQT*IWSWTLFWLVSY
6130	36498	A	6179	1	2022	
6131	36499	Α	6180	1	882	
6132	36500	C	6181	1	1935	
6133	36501	A	6182	1	672	
6134	36502	В	6183	140	2195	
6135	36503	Α	6184	26	354	

NO.		IVECT	ISEC ID NO.	Nucleotide	Muciconde tocation or last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	<u> </u>	<u> </u>				
6136	36504	Α	6185	86	747	SLEDAMSLKDDDSGDHDQNEE
		ł				NSTQKDGEKEKTERDKNQSSS
					1	KRKAVVPGPAEHPLQYNYTFW
		ļ				YSRRTPGRPTSSQSYEQNIKQIG
		ŀ				TFAS\VEQFWRFYSHMVRPGDL
						TGHSDFHLFKEGIKPMWE\DDA
						NKNG\GKWIIRL\RKGFGSRC\W
						ENLILAMLGEQF\MVGEEICGG
		l				CGVCPAFQEDIIS\IWNKTASDQ
						ATTARIRDTL\RRVL\NLPANTI
						MEYKN
6137	36505	A	6186	23	127	
6138	36506	A	6187	2	419	
6139 6140	36507	A	6188	19	714 373	AVFFFFFFFFFFFFFFFFFFFFFFFFFFFFFFFFFFFFF
6140 36508	Α	0189	119	3/3	FFFFFFFFFFFSSSSSSFFFFSFFL	
		l				FFFFKLKELRLVI*PLNPEN*ESC
			1			PLFSPVNLDMP*TFKDKRWKV
						NWARSTSRQQLLESARTKSWA
(1.41)	12.6.500	 _	(100	227	120	EREGGTRWRAW KNFLPNCSAKELFFFFFFFFFFL
6141 3	36509	A	6190	227	420	i
						LLLLLLYFKF/MI*WFYKHLAF
6142	26510	ļ.	(101	400	582	PLLAFILSPTAL*RGVFHH
6142	36510	A	6191 6192	1202	1678	
6143	36511	A A	6193	2	109	FTFWHDFAAAGTGCSFPCLVLP
6144	36512	A	0193	2	109	SWW*QNLSAFACL
6145	36513	Α	6194	110	377	QWISRQKPYKPEKRGDQYSTFL
0143	30313	A	0194	110	377	*EC*ILVPPLFWLVGFLPRDPLL
	ŀ					V*WAYGCP*HFFLHFNLGESNN
			•			YVSWGCSS
6146	36514	1	6195	1	381	1 v 3 w 0 c 3 5
6147	36515	A	6196	1	759	
6148	36516	A	6197	1	609	
6149	36517	A	6198	2	1178	
6150	36518	В	6199	159	290	
6151	36519	A	6200	716	1001	SMPASTHLILVGSCSWKMVIVL
0131	30319	^	0200	1,10	1001	PLVTHFLFTALTVPWNLPWVEL
		ŀ				YWNM*TM*LRSMKESLMATIF
						TLPELKA\VLHEKLQ*LQSTEDL
			:		•	*HGLLGG
6152	36520	A	6201	5772	5971	TANIIMTGSNSHITILTLNVNRL
0132	130320		0201	3112	37/1	NARIKRHRLENCIKSQDPSVCCI
		1				*ETHLMCRDTYRLKIKGWSN
6153	36521	A	6202	1	974	E HENCKET I REKIKG W SIV
10177	36522	A	6203	1	600	

NO:				Nucleotide		Amino acid sequence (X=Unknown,
1	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
1	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
6155	36523	Α	6204	51	424	MAPKQDPKPKFQEGERVLCFH
l						GPLLYEAKRVKVAIKD/ITREI/H
1		ŀ				SYITVVGIKIGMNGFRRAKYSN
ĺ			<u> </u>			TWTPICRNSENFKKPISKFVL*T
ĺ						EY*GEMPVD*FWTWNEQWRSC
ĺ						WLPCLYNDTSDLSAGV
6156	36524	Α	6205	1	3827	MQKAIRLNDGHVAPLGLLARK
						DGTRKGYLSKRSSDNTK WQTK
						WFALLQNLLFYFESDSSSRPSG
						LYLLEGCVCDRAPSPKPALSAK
						EPLEKQHYFTVNFSHENQKALE
						LRTEDAKDCDEWVAAIAHASY
						RTLATEHEALMQKYLHLLQIVE
	ļ					TEKTVAKQLRQQIEDGEIEIERL
						KAEITSLLKDNERIQSTQTVAPN
	ļ					DEDSDIKKIKKVQSFLRGWLCR
						RKWKTIIQDYIRSPHADSMRKR
						NQVVFSMLEAEAEY
6157	36525	A	6206	1	900	
6158	36526	С	6207	172	501	
6159 36	36527	Α	6208	499	848	KIRVPAAVAGLSPPLGPWSLRS
						PSQPERCCPWRNSSWWTAPRTS
						IITAAKGVSPARLSSISCTTRGS
						WVKTPTPTRARMVIASSNLERP
						SALSRM*PTSQSMTRKRWWRL
		ļ				WPSTTL
6160	36528	A	6209	3	495	SCAQTFPKLDTFLEHIKSHQEEL
		1				SYRCHLCGKDFPSLYDLGVHQ
						YSHSLLPQHSPKKDNA/CLQVF
						PCERYLRRHLPTHGSGGRFKCQ
						VCKKFFRREHYLKLHAHIHSGE
		l				KPYKCSVCESAFNRKDKLKRH
						MLIQYDEVRQGLGMVPGTQKV LNKCFCHNVKKKK
6161	26520	_	6210	667	2142	LNKCFCHNVKKK
6161 6162	36529 36530	A	6211	667	396	
6163	36531	A	6212	213	446	
6164	36532	A	6213	101	580	VAGVGSLPRSMEDDAPVIYGLE
104	30332	 	213			FQARA\LTPQT\AETDAIRFLVG
						TOSLKYDNOIHIIDFDDENNIIN
1						KNVLLHQS/AGEIWHISASPADR
1						GVLTTCYNRTSDSKVLTCAAV
1						WRMPKELESGSHESPDDSSSTA
1						QTL\ELLCHLDNTAHGNMACV
						VWEPMGDG
	36533	Α	6214	1	933	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
6166	36534	A	6215	3	404	RLKEKKLVKEVIAVSCGPAQCQ ETIRTALAMGADRGIHVEVPPA EAERLGPLQVARV/AVVTADLR LNEPCYATLPNIMKAKKKKIEV IKPGDLGVDLTSKLSVISVEDPP QRTAGVKVETTEDLVAKLKEIG
6167	36535	Α	6216	1	408	
6168	36536	A	6217	88	871	MADVRVLVAVKRVIDYAVKIR VKPDRTGVVTDGVKHSMNPFC EIAVEEAVRLKEKKLVKEVIAV\ SCGPAQCQETI\RTALA\MGA\D RGIHVE\VPPAEAERF/GVPLQV ARVLGQLAKK\EKVDLVLLGK QA\IACTDCNQ\TGQMTAGFLD WPQGTF\ASQVTLEGDKLKVER EIDGGLETLRLKLPAVVTADLR LNEPRYA\TLP\NIMKAKKKIE VIKPGDLGVDLTSKLSVISVEDP PQRTAGVKVETTEDLVAKLKEI
6169	36537	Α	6218	1	1023	
6170	36538	Α	6219	110	842	
6171	36539	Α	6220	1	122	
6172	36540	A	6222	727	941	MNDATA VQEGSQMYSWQKLT PTGPFIMQPEGMGRLFAINKMA EGPFPEHYEPIETPLGTNPLHPN VVSNPVVRLYEQDALRMGKKE QFPYVGTTYRLTEHFHTWTKH ALLNAIAQPEQFVEISETLAAAK GINNGDRVTVSSKRGFIRAVAV VTRRLKPLNVNGQQVETVGIPI HWGFEGVARKGYIANTLTPNV GDANSQTPEYKAFL/GQHREGV RN*LHHAAFSGA*LQCFQYAH NSRKNN WLRMMEHRILFPLWKRESKKA
6173	36541	A	6223	737		HYVQSSVLLCLGMLR*KYKGK KGKKERKKKDEKRKGNGKKRI KIHSI
6174	36542	Α	6224	1	1072	
6175	36543	A	6225	1	585	
6176	36544	A	6226	100	930	A OFFUNI OF AN ANGO COLORS
6177	36545	A	6227	28	392	AGEVPLÖLPPAAVCSACLORPA LWWNSSFFL*SCLHALFH*VDL QSLIAFLLLD*SSY*YLCMLHEV LVLCFSAPSGHLCSSLS*LF*LAI HLTFFQGS*LPCIGLEHASLAQR SLLLPTF
6178	36546	A	6228	1	271	SAIVTFVCFMKFSCCVFQLHQV FHVL\FKLVILVSSSCNLFSRFLA SLHWVTTCSFSSEEFVITHLLKP APVNSSNSFSVQFCSLAGKEL

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO:	Nucleotide location of first	•	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO:	sequence	noa	09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
6179	36547	I A	6229	11	1089	MGWRPRWLPYLYWGFQSDLC
		-				PHCKRKAQEEDSLAAQAASHP
	ļ					RLSANSKNNSWGKQYSYALFK
						AMRHMLCIGYGRQAPAAVSDV
						WLTMLSM\IVGATCYAMFIGH/
						ATALIQVLDCSRRQYQEKVVLP
						REHPNIGSSQEGGTWHQTVGL
						QQTSDKFDANDPILKDQTQEW
						SGSATFTSDGKIRLFYTDYSGK
						HYGKQSLTTSAANTGTENGYQ
						GEESLFNKAYYGGGTNFFRKES
						QKLQQSATTRDAELANGALGII
						ELDNDSPLKKRGATQRSSEENI
						TLKAYKRDVRSLLMLTRNEDV
						ADSLNSSKKRQNYQLPSIRSVT
İ		1				RCKKIGVCAKGIGCGQGTHYLS
					}	VSILNTADRDRRKLGFTRTSMY
6180	36548	D	6230	1	1140	VSILNIADRDRRRLOFIRISMI
6181	36549	B A	6231	3	204	GSDYELGLW*PWLTSEATDLH
0101	36349	A	0231	3	204	GECYSS*SGASGVVRSSLWARG
						LAGFRSEAADLRGRESVAVTN
6182	36550	-	6232	1	1296	LAUFKSEAADLKUKESVAVIN
6183	36551	A B	6233	1	1686	
6184	36552	A	6234	1	627	
6185	36553	A	6235	570	736	SATFGRSSRSGCSG\FPAASRRG
0185	30333	^	0233	370	730	RLPLSPPPWPAV\RVYPCGMLD
		1				NKASAAHPPKKK
6186	36554	A	6236	1	103	NKASAATITKKK
6187	36555	A	6237	699	887	GLQPPAPRETRPLY*TLAWNLH
0.07	30333	`	10237			FLFVSPLDSCDLVRSVLTCLSPP
1		l				GNSPSFDMHRYQLPCVH
6188	36556	A	6238	2	803	C1181818181
6189	36557	A	6239	2	1277	YKETYLIHLFHTFTGLSIAYFNF
						GNOLYHSLLCIVLOFLILRLMG
						RTITAVLTTFCFQMAYLLAGYY
						YTATGNYDIKWTMPHCVLTLK
						LIGLAVDYFDGGKDQNSLSSEQ
İ						QKYAIRGVPSLLEVAGFSYFYG
						AFLVGPQFSMNHYMKLVQGEL
1						IDIPGKIPNSIIPALKRLSLGLFYL
						VGYTLLSPHITEDYLLTEDYDN
						HPFWFRCMYMLIWGKFVLYKY
		1				VTCWLVTEGVCILTGLGFNGFE
						EKGKAKWDACANMKVWLFET
						NPRFTGTIASFNINTNAWVARYI
						FKRLKFLGNKELSQGLS\MLFLF
		1				_
		1				LWHGLHSGYLVCFQMEFLIVIV
						ERQAARLIQESPTLSKLAAITVL
						QPFYYLVQQTIHWLFMGYSMT
						AFCLFTW\DKWLKVYKSIYFLG
	<u>i</u>		<u> </u>	<u> </u>	L	HIFFLSLLFILPYIHK\AMVPRKE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
6190	36558	I _A	6240	56	324	
6191	36559	A	6241	68	487	
6192	36560	Α	6242	52	457	
6193	36561	A	6243	263	787	SLEV*KWPGVACRLQL*P/MVP
						APKS*ESKSAGSLNTPSAICRDA SGRGLARPWLRKSKCLACHLV DTRSKTAGGLAPVPSAHLISVR EREPGLCAVLVSQLEFRVGVGL VGPTLGARREPCWPQAMGDLA PEPVAALRLISRRALPAFPWGR ARDLQPAMPEPPTHSMGSCAPE PPR
6194	36562	Α	6244	3	236	
6195	36563	С	6245	1	1608	
6196	36564	В	6246	1	1083	
6197	36565	A	6247	823	1055	KQKDGTGAPWSRGRRSSGRLG THRSPRRGWEAQAWRAAGPET CPAGRQLRPGEKSSAAPG*RSR RKDAYEMDTAKKK
6198	36566	В	6248	1	390	
6199	36567	Α	6249	27	363	
6200	36568	В	6250	1132	1268	
6201	36569	Α	6251	11	489	
6202	36570	Α	6252	356	638	
6203	36571	Α	6253	I	507	
6204	36572	Α	6254	80	336	
6205	36573	В	6255	100	912	
6206	36574	В	6256	821	2992	
6207	36575	Α	6257	2	602	
6208	36576	Α	6258	1	519	
6209	36577	A	6259	3	228	STGVPVSQ\RTCLLEKASVLFNT GALYTQIGTRCDRQTQAGLESA IDAFQRAADMSPPGLTRRSLGP TRWHQGTPH
6210	36578	Α	6260	1	963	
6211	36579	Α	6261	1	925	
6212	36580	A	6262	1	397	
6213	36581	Α	6263	1	2625	
6214	36582	A	6264	11	241	SLQHLNPATFLPVSKSPVKHNC VEVLDSVYSSGPNHRDHP*TSV DWELYMDGSSFTNPCKVTLKK TTSPAPVTPGS
6215	36583	Α	6265	1	1662	
6216	36584	В	6266	1	897	
6217	36585	Α	6267	3	258	
6218	36586	Α	6268	1	279	
6219	36587	Α	6269	1	258	
6220	36588	Α	6270	1	639	
6221	36589	В	6271	1	1935	
6222	36590	Α	6272	698	1643	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
6223	36591	A	6273	91	386	IFDFLFLLYVSGCNSIAEPQHFT TTVTRCSPTVAFVEFPSSPQLKN DVSEEKDQK\KP\ENEMSGKGG SWVLFTKGEVQEWRESGLNPT KFNMGPLIYS
6224	36592	Ā	6274	1	2538	
6225	36593	Α	6275	1	471	
6226	36594	A	6276	152	520	CPLTLLLPSRGAGLGTCSPPCLS LPPTPWAPVRPEPPRRAPPPAPR RPVPSTGKEQTTQGLRNASARR GTGRQLHL\SPGVGSTG*SQGL
6227	36595	A	6277	831	1449	HPGQWLRRVCWVSSGAGPLAL CSISCPALAAFPRVGLGTCSPPC LSLPPTPWAPVRPEPPRRAPPPA PR/PPSPIDHPRAEEYE/PHGAGL AGSSTCSPSAESTG*SQLGS*VV RVRGFILEVSETKNPPIPDTFGL PPGSSSAASTLVQVTRL*ISAISL PHLLSRPCSNTARGPVPAPVPG AAVRAPAAGGRWCGGARGRG RRGGGH
6228	36596	C	6278	1	426	
6229	36597	A	6279		535	MKHSGAQLASPSGS\PLGPQVE LPASPVPCACTSQPLGGRWDW APWSRGWCSSGRLGPHRSPWS GWE\LRHGGLQVPSPAPREGS* GPARNP/GTAPGRPAAPS\GGPP SPRPPRTPAGPQALRAAPVP\LA PLPPHLPAS*GSGLQPWPA*KE APTVQCQAEGLLKCRQSGSPGR GGAESE
6230	36598	В	6280	1	1743	
6231	36599	Α	6281		428	GTREEHLKRSGREIALPIEACV MLLLETGMKEEGLFRIGAGASK LKKLKAALDCS\ILTWDEFYSD PHAVAG\ALKSYLRELPEPLMT FNLYEEW\AQVASVQDQGQKT FKDLWRTCQKLPPQNFVNFRY LIKFLAKLAQTSDV
6232	36600	A	6282	135	512	LPFRGAGLETCSPPCLSLPPPPW APVRPKPPR*APPPAPRRPVPST TQGSLGPHCFYEL*HSP*RSAAS LLKPARP*AHREERTTP/GRSAL/ TSCNTHREGLQ/PSLRKKETPNT SEHQKEQTPDAPP

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
6233	36601	A	6283	698	2233	IQAAMQAAPQFGPSTED*CCPIT
				-		SEAPWAIMDAKLR\LKPPSHPPL
						VSPHLNPQVWDTSSPSLATEHA
						SLTISLKPNHPYPAQSQYSIPQH
						ALKGSKPVITRLLEHGLLKPINS
						PYNSPILPVQKLDKSYRLVQHL
	1					CLIYQIVLPIHPMVPNPYTLLSSI
	ŀ					PFTTHYSVLDLKHAFFTIPLYPS
	ļ	l				SQPLFAFTWTDPDTHQAQQITL
						AVLPQCFIDSPHYFSQAQISSSS
						VTYLGIILMKTHIGLGAVEQGV
						VLVGEARAAQEPMEWVGGSG
		1				MVGCRSRALPRGKAAKARREI
		ŀ		İ		ERSAVTIVPVLDFNPAFHIIPDTT
:	ļ					PDHHDCISLIHLTFTPFPHISFFP
						VPHLEHTWFIDGSSTRPNCHSP
						AKAGYAIVSSTSIIEATALPRST
						TSQQAKLIAYTDSKCAFHILHH
		ŀ				HAVIWEERNFLTMQGASIINAF
						LIKTLLKDILLPKEAGVIHCKGH
	İ					
						QKASDPITQDNAYADKVAKKQ
						LAFQLLSLTAVFLLLIRPLSPTSP LKLPLINLFPHKANDS
6234	36602	A	6284	1	1239	LKLPLINLFPHKANDS
6235	36603	A	6285	1	477	QRPPHGVFVLGVAGRVHGPEL
0233	30003	A	A 16283	1	777	ASHKAGPQLGDVGVCAGQGG
						QVGPLVLQPVVEHIQQLPG/AG
	ļ					
	ļ	l				HCAHQ*QGSPS*SSLPRPERSLQ
		1				CRRPQGATATIRSTWWGPRGP
		1		ļ		VCTAALVPLCWPGWPVQKDEL
						QAVRSHSALASAMQTCSLAGP
6236	36604	A	6286	1	1134	GLGSSHARLTA
6237	36605	A	6287	673	836	
6238	36606	A	6288	1	225	
6239	36607	Α	6289	1	468	
6240	36608	Α	6290	495	1637	ESQVLSNVSPHVKV*NMASWE
						GKDLTVPQPDTRKGSVLRRISK
						RGRNASCS*DKRKASVSCLSLG
]				1		NGMSRYKTRLYAPSTEIGKNRL
						RAGGGTCGQQYCFVKH
6241	36609	Α	6291	1	507	
6242	36610	Α	6292	215	389	
6243	36611	A	6293	1	255	
6244	36612	В	6294	69	1055	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
6245	36613	A	6295	23	1596	IKPQRWGETRAEMLKILKTREP FLLQRIASPRQQWNKAGQRRS WFFERL/NKIDRLLARLIKKKRE KNQIDAIKNDKGDITTDPTEIQT TIREYYKHLYANKLENLEEMD KFLDTYTLPRLNQEELESLNRPI TGSEIEAIISSLRTKKSPGPDGFT AEFYQRYKEELVLFLLKLFQSIE KEGILPNSFYEASIILIPKPGRDT TKKENFRLISLMNIDVKILNKIL ANRIQQHIKKLVHHDQVGFIPG MPDWFNICKSINIIHHINRTNGK NHMIISIDAEKAFEKIQQCFMLK TLNKLGIDGMYLKIIRAIFDKPT ANVILNGQKLEAFPLKTGTTQG CPLSPLLFKIVLEVLARAIRQEK EIKRIQLGKEEAKLSLFADDMIV YLENPIVSAQNLLQLRSNFSKV LGYKINMQKSQAFLYTKNKQT ESQIMSELPFTIATKRIKYLGIQL TRDVKSLFKENYKPLLSKIKED TNKWKNIPSSWIGRISIAKMAIL
6246	36614		6296	1	1488	PKVIYTFNAIPIKLPMTFFTELEK
	1	A				
6247	36615	A	6297	1	1401	
6248	36616	Α	6298	2	1256	
6249	36617	В	6299	84	955	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
6250	36618	Α	6300	735	2718	RSFPRSRPSPFLLLSRYLRIHMV
		1				FSVLPFGLQNPKYLLSGSLQEK
		İ				FRTPGINSHKTLDPNRVIIKVRR
	İ					RLEKEKALRAYVGKSEVRNMH
		1				LRERCKL*KKREKNQIDTIKND
		ļ				KGDITTDATEIQTTIREYYKHLY
						ANKLENLEEMDKFLDTYILPRL
						NQEEVESLNRPITGSEIEAIINSL
						PTKRSPGPDGFTAEFYQTYKEE
		1				LVPFPLKLFQSTEKEGILPNSFY
				•		EASIILIPKPGRDTTKKENFRPIS
						LMNIDAKILNKILANRIQQHIKK
						LIHYDQVGFIPGMQGWFNKRK
	İ					SINVIQHINRTNDKNHMIISIDAE
						KAFDKIQQPFMLKTLNKLGIDG
						TYLKVIRAIYDKPTANIILNGQK
						LEAFPLKTDTRQGCPLSPLLFNI
						VLEVLARAVRQEKEIEGIQLGK
	1					EEVKLSLFADDMIVYLENPIVS
						AQNLLKLISNFSKVSGYKINIQK
	-					SQAFLYTNNRQTESQIMSELPFT
						TASKKIKYLGIQLTRDVKDLFK
						ENYKPLLNEIKEDTNKWKNIPC
						SWVGRINIMKMAILPKVICRFN
						AIPIKLPMPFFTELETTTLKFIWN
						QKRARKSILSQKNKAGGITLPD
						FKLYYKATVTKTAWYWYQNR
						DIDQWNRTEPSEIMPHIYNYLIF
						DKPEKNKQWGKDSLFNK*CWE
		ŀ				NWLAIWRKLK
6251	36619	В	6301	1	2907	
6252	36620	Α	6302	1	2358	
6253	36621	A	6303	1	1866	
6254	36622	Α	6304	1	885	WO DISCOURANT LINE OF COOK
6255	36623	Α	6305	1	1059	MKNNSESFEPLLLIPRQRGSGV
						DLQQTPTDLQLRVLTVRRKTN
		1				KQDIHTKTPTVRHHHQRPKNP
						WDTFKAVCRGKFIALNAHKRK
						QERSKIDTLTSQLKEVEKQEQT
		1				QSKASRRQEITKIRGELKEIETQ
	1					KTLQKINESRSWFFEKINKIDRL
						LARLIKKKREKNQIDTIKNDKE
						DITTDSTEIQTTIREYYKHLYAN
	1	1				KLENLEEMDKFLNIYTLPRLNQ
						EEVESLNRPITGSEIEAIINSLPT
		1				KKSPGPDGFTAEFYQSYQEELV
		1				PFLLKLFQSIEKEGILSNSFYKA
						NIILIPKPGRDTTKKENFRPISLM
		1				NIDAKILNKIL/ANRIQQHIKKLI
						HHDQVGFIPGMQGWFNI
6256	36624	Α	6306	1	1692	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
6257	36625	В	6307	1	2871	
6258	36626	Α	6308	2	2315	
6259	36627	Α	6309	1	1210	MMSFHELSVHTAIDGHSHSSCS
i		ŀ				AWCSPIDRDYFEQENNSSSSCC
		ŀ				CCCYHCHRKHCGGKGLSFPGE
						MPLSPGHFNKMMYSVSELMQG
						SGSGTFIPLPAELSLWLLVTNTK
		l				GQRMIKNDAEIQTTIREYYKHL
						YANKLENLEEMDKFLDTYTLP
						RLNQEEVESLNRPITSSEIVAIIN
						SLPTKKSPGPHGFTAEFYQRYK
						EELVPFLLKLFQSIEKEGILPNSF
						YEASIILIPKPGRDTTKKENFRPI
						SLMNSDAKILNKILANRIQQHIK
						KLIHHDQVGFIPGMQGWFNICK
						SINVIQHINRTKDKNHMIISIDAE
		1		}		KAFDKIQQPFMLKTLNKL\GIK
						YLGIQLTWDVKDLFKENYKPL
						LKEIKEDTNKWKNIPCSWVGKI
						NIMKMAIPPKATSPLESYKREK
6260	36628	Α	6310	1	2745	MEGEMNEMKREGKFREKRIKR
						NEQSLQEIWDYVKRPNLRLIGV
		ŀ				PESDGENGTKLENTLQDIIQENF
						PNLARQANVQIQEIQRTPQRYS
						SRRATPRHIIVRFTKVEMKEKM
		İ				LRAAREKDRSTRQKVNKDTQE
						LNSALHQADLIDIYRTLHLKSTE
						YTFFSAPHHTYSKIDHILGSKAL
						LSKCKRTEIITNYLSDHSAIKLE
						LRIKNLTQNRSTTWKLNNLLLN
						DYWVHNKMKAEIKMFFETNEN
						KDTTYQNLWDTF

SEQ ID NO:			SEQ ID NO:		Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
6261	36629	A	6311	1	2052	MGDFNTPLSTLDRSMRQKVNK
						DTQELNSALHQADLIDIYRTLH
						PKSTEYTFFSAPHHTYSKIDHIV
						GSKALLSKCKRTEIITNYLSDHS
						AIKLELRIKNLTQNRSTTWKLN
						NLLLNDYWVHNEMKAEIKMFF
						ETNENKDKTYQNLWDAFNAV
		ŀ				CRGKFVALNAHKRKQEGSKID
						TLTSQLKELEKQEQTHSKASRR
						QEITKVRAELKEIETQKTLQKIN
		ŀ				ESRS\ARLIKKKREKNQI/DAIKN
	}					DKRDITIDPTEIQTTIREYYKHL
		1			·	YANKLENLEEMDKFLDTYTLP
						RLNQEEVESLNRPITGSEIVAIIN
						SLPTKKSPGPEGFTAEFYQRYK
				i		EELVPFLLKLFQSTEKEGILPNS
j						FYEASIILIPKPGRDTTKKENFRP
		İ				ISLMNIDAKILNKILAKRIQQHI
		l				QKLIHHDQVGFIPGMQGWFNIC
						KSINVIQHINRTKDKNHMIISID
•						AEKVFDKIQQRFMLKTLNKLV
						LEVLARAVRQEKKIKGIQLGKE
						ELKLSLFADDMIVYLENPIVSA
						QNLLKLISNFSKVSEYKINVQKS
						QAFLYTNNRQTESQIMSELPFTI
	1					ASKRIKYLGIQLTRDVKDLFKE
						NYKPLLKEIKEDVNKWKNIPCS
						WVGRINIVKMAILPKVIYRFNAI
						PIKLPMTSFTELEKTTFKFIWNH
						KGARIAKSILSQKNKAGGITLPD
						FKLYYKATVTKTAWY

SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			sequence		
36630	Α	6312	2	2143	SSDGSWWTGFQWREWRQAGR
}	ļ				SVNSWDNPKQEVRASSKDKSR
					GSIQEAMRMQSSAKLLCSAWT
					LAYSIAVRTLSSDSEGQPPLVIH
		:			RQTGSGEDLQQTPTDLQLRVLT
1				•	IRRKTNKQKGHPHQNPISSRRQ
					EITKIRAELKKIETQKPFKKINES
					RSWFFEKINKIDRLLARLIKKKI
					EKNQIDAIKNDKGNITTNPTEIQ
1					TTIREYYKHLYANKLEHLEEM
					DKFLDIYTLPRLNQEEVESVNR
					PITGSEIEAIINSLPTKKSPGPDRF
					TAELYQRYKEELVPFLLKLFQSI
					EKEGILPNSFYEASIILISKPGRD
					TTKKENFRPISLMNIDAKILNKI
					LANQIQQHIKKLIHHHQVGFIPG
					MQGWFNILKSINVIHHINRTKD
					KNHMIISIEAEKAFDKIQQPFML
					KTLNKLGIDGTYLTYLKIIRAIY
					DKPTANIILNGQKLEAFPFKTGT
					RQGCPLSPLLFNIVLEALARAIR
					QEKEIKGIQLGKEDVKLSLFAD
					DMIVYLENPIVSAQNLLKLISNF
					SEVSGYKINVQKSQAFLYTNNR
					QTESQIMSELPFTIASKRIKYLGI
					QLTRDVKDLFKENYKPLLNEIK
					EDTNKWKNIPCSWIGRINIVKM
					AILPKTLNQKFSYWFRVNKHYI
					HQRTFPLKETEFNTIATLYNGA
					SP/CTAPKSTGTNGHQASGLPRF
					*RIAFCSALVKSKRKLYQGYLP
					GQTDRREEGVSWCPGGP
36631	Α	6313	1	3018	
36632	A	6314	1	2016	
36633	В	6315	1	1215	
	36631 36632	36631 A 36632 A	of peptide sequence hod in USSN 09/540,217 36630 A 6312 36631 A 6313 36632 A 6314	36631 A 6313 1 36632 A 6314 1	1

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
6266	36634	Α	6316		2268	MAGYPSETKPPEERSDSNICSSA
						IFTVLHPPLLIPRQTAFGMDLQQ
						MPTDLQLRVLTVKRKTNKQKG
		:				HPHQNPIRTSPSSKTEDFQPTKI
		į				KRDKEGHYIMVKGSIQQEELTI
						LNIYAPNTGAPRFIKEVLRDLQ
						RDLDSHTIIMGEFNTRLSTLDRS
						MRQKVNKDIQELNSALHQADL
						IDISRNLHPKSTEYTFFSAPHRT
						YSKIDHIVGSKALLSKCKRTEII
						TKCLSDHSAIKLELRIKNLTQNC
						STTWKLNNLLLNDYWVHNEM
						KAEIKMFFETSENKDTTYQNL
						WDTFKAVCRGKFIALNAHKRK
						QERSKTDTLTSQLKELEKQEQT
				ļ		HSKASRRQEITKIRAELQEIETQ
]	ļ		RTLQKISESRSWFFEKINKIDRS
						LARLIKKKREKNQIEAIKNDKG
			•			DITTNPTETQTTIKEYYKHLYK
		l		į		NKLENLEEMDKFLNTYTLPRLN
		ł				QEEVESLNRPITGSEIVAIINSLP
		1				TKKSPGPDGFTAEFYQRCKEEL
						VTFLLKLFQSIEKEGILPNSFYE
						ASINLIPKPGRDTTKKENFRPISL
		l				MNIDAKILNKILANRI/WGN*AE
						ERNKEYSIRKRGSQIVPVCR*H
						DCVSRKPHHLSPQSP*ADKQLQ
			•			QSLRIQNQCTKITSILIHQ*QTNR
						EPNHE*TPIHNCFKENKIPRNPT
						YKGCEGPLQGELQTTAQ*NKR
						GHKQMEEHSMLMDRKNQHRE
						NRHTAQGNL*IQCHSHQATNDF
6267	36635	A	6317	1	1797	
6268	36636	Α	6318	2	2063	
6269	36637	Α	6319		1212	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	1			sequence		:
6270	36638	A	6320	1	[2718	MALFRVGMTTRRLGHANKFTR
0270	30030	 	0320		27,0	DLLVKRLLLYEFVVEIIPGLRTK
			<u> </u>			TTLVGPLETGYTSSDSVNSPHF
						MLDIALSHNVKSTQLLLRPWR
						NIDCSRNHKNAKKEEQMGDDE
						INRQQECSSSPAMEQSWTENDF
						DELREEVFRRSNYSELQEEIRTN
						GKEVKSFEKKLDEWITRITNAE
		ļ				KSLKHLTELKTKARELQLEKQE
!						LTHSKASRRQEITKIRAELKEIE
						TQKTLQKINESRSWFFEKINKID
						RPLARLIKKKREKNQTDTIKND
			:			KGDIITDPTEIQTTIREYYKHLY
						TNKLENLEEMDKFLDTYVLPRL
						NOEEVESLNRPITGSEIEAIINSL
						PTKKSPGPDGFTAEFYQRYKEE
		į				LVPFLLKLSQSIEKERILPNSSYE
						ASIILIPKPGRDTTKKENFRPISL
			i I			MNIDAKILNKILANRTQQHIKK
						LIHHEQVGCIPGMQGWFNIRKS
						INVIQHINRTKDKNHMIISIDAE
						KLISKFSKVSGHKINVQKSQAF
						LYTDNRQTESQIMSELPFTIASK
						RIKYLGIQLTRDVKDLFKENYK
						PLLNEIKEDTNKWKNIPCSWVG
			:			RINIVKMALLP\RFSAIPIKLPMT
						FFTELEKTTLKFIWNQKEP/CIA
						KSFLSQKNKAGGITLPDFKLYY
						KATVPKTAWYCYQNRDIDQW
			:			NRTEPSEIMLLIYNYLIFDKPDK
						KKEWGKDSLFNKWCWENWLA
			1			ICRKLKLDPFLTPYTKMNSRWI
6271	36639	A	6321	1	2307	
6272	36640	Α	6322	1	1989	
6273	36641	Α	6323	1	3285	
6274	36642	A	6324	3	4732	
6275	36643	В	6325	1	2693	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	l	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
6276	36644	Α	6326	603	2976	DNTARGTIRQQHLLFTNIRCSA
		İ				ASAADTQANRVWSGPPANSNR
		İ				PAA\RVLTVRRKTNKQKGHPH
1		İ				QNPICTSPSSKTKETQTTIREYY
		Ì				KHLYANRLENLEEMDKFLDTY
						TLPRLNQEEVESLNRRITESDIE
						AIINSLPTKKSPGPDGFTAEFYQ
						RYKEELVPFLLKLFQSIEKEGIL
						PNSFYEASIILIPKLGRDTTKKE
						NFRPISLMNIDAKILNKILANRI
	·					QQHIKKLIHHDQVGFIPGMQG
						WFNICKSINVIHHMKRTKDKNH
						MIISIDAEKAFDKIQQPFMLKTL
						NKLGIDGTYLKIIRAIYDKPTAN
						IILNGQKLEAFPLKTGTRQGCPL
						SPLLFNIVLEVLARAIRQEKEIK
						GFQLGKEEVKLSLFVDEMILSL
1						ENPMVSAKNLLKLISNFSKVSG
1						YKINVQKSQAFLYTSNRQTESQ
						IMSELPFTIASKRIKYLGIQLTGD
						VKDFFKENYKPLLNEIKEDTDK
				ł		WKNIPCSRVGRINIMKMAILPK
						VIYRFNAIPIKLPMTFFTELEKTT
1		ļ				LKFIWNQKRARIAKSILSQKNK
						AGGITLPDFKLYYKATVTKTAC
						HRVGRAQQHSISKWKWYIHDW
		ŀ	}			SQVGPEGTNDSARYPDTTQKW
1						TAAALQPLSRTSLKDSHEGKSS
			-			QWAELRAVHLVLRFAWKEKW
						PDVQLYTDSWAVASGLAGWS
						GTWKKHDWKIGDKEIWGRATP
			İ			VIAQWAHEQRGHGGRDGDYA
6277	36645	Α	6327	3193	6107	SMKTGLEKKMKRNEQSLQEIW
						DYVKRPNLPLIGVPESDGDNAT
						KLENTLQDIIQENCPNLARQANI
						QIQEIQRTPQRYSSRRATPRHIIV
						RFTKVEMKEKMLRAAREKGRV
						THKGKPIRLTAALSAETLQARR
						DRRPIFNILKEKNFQPRISYPAK
						LSFVSEGEIEYFTDKQMLRDFL
						MTRPALKELLKEALNMETTGA
			:			PRFIKQVLRELQRDLDTHTIITG
						DFNTPLSTSDRSTRQKVNKDIQ
						ELNSALHQADL
6278	36646	Α	6328	1	1808	
6279	36647	Α	6329	1	4695	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence		in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
		<u> </u>				
6280	36648	A	6330		2915	MIMGDFNTPLSTLDRSTRQKVN KDTWELNSALHQADLIDIYRTP HPKSTEYTFFLAPHHTYSKIDHI AGNKALLSKCERTEIITNCLSDH
						SAIKLQLRIKKLTQNHSTTWKL NNLLLNDYWVHNEMKAEIKIF FETNKNKDTTYQNLWDTFKAV CRGKFIALNAHKRKQERSKTDT
						RTSQLKHLENQEQTHSKASRRS WFFEKINKIHRPLARLIKKKREK NQIDAIKNDKGDITTNPTEIQTTI
					:	REYYKHLYTN
6281	36649	В	6331	1	6730	
6282	36650	Α	6332	1	378	
6283	36651	Α	6333	3	1700	
6284	36652	Α	6334	1	1759	
6285	36653	Α	6335	2	409	
6286	36654	A	6336		2051	MFGRSRSWVGGGHGKTSRNIH SLDHLKYLYHVLTKNTTVTEQ NRNLLVETIRSITEILIWGDQND
						SSVFDFFLEKNMFVFFLNILRQK
		1				SGRYVCVQLLQTLNILFENISHE
						TSLYYLLSNNYVNSIIVHKFDFS DEEIMAYYISFLKTLSLKLNNH
						TVHFFYNEHTNDFALYTEAIKF
						FNHPESMVRIAVRTITLNVYKV
						DNQAMLHYIRDKTAVPYFSNL
						VWFIGSHVIELDDCVQTDEEHR
						NRGKLSDLVAEHLDHLHYLND ILIINCEFLNDVLTDHLLNRLFLP
						LYVYSLENQDKVFLIIHHAPLV
						NSLAEVILNGDLSEMYAKTEOD
					:	IQRSSVLPTLSSLWQGSHLSLNQ
						LQSGLHKCSSHLCGAQAAADS VTGEIPAIRSLEWLISAGSKART
						FFFLKMLIGFWEKVDCEYQRR
		1				QVLSTRLQEALPSNRLTDVAAV
						HSSCMLGFGSTAPRGSWIGDPA
		1				AVHLPLPGELAEHLGSKGTTTV
						TKHQPQAKPSIRCFIKPTETLER
		1				SLEMNKHKGKRRVQKRPNYK
						NVGEEEDEEKGPTEDAQEDAE
						KAKGTEGGSKGIKTSGESEEIE
						MVIMERSKLSELAASTSVQEQN
						TTDEEKSAAATCSESTQWSRPF
						LDMVYHALDSPDDDYHALFVL
			}			CLLYAMSHNK/GKSPEKEEGLS
6007	26655	1	6227		2752	GTQSHPGKAGTFGKEGAEERK
6287	36655 36656	A	6337 6338	2	2753 577	
6288	<u> </u>	-	Ļ	1	849	
6289	36657	Α	6339		047	<u> </u>

possible nucleotide ble nucleotide insertion)
,
OACAEL CREWORK
QASAFLGRTWSRP
VSCLAKAGVQGYL
RFLFSERTGSFSKL
GLGSGLGQSLGQV
QISNFTKDMLMEG
DSRTKEIEAIHAIL
CLCTDLEEKHEASE
/RNQLQQKEVEISH
DQFLKLQSAAQSV
TASSSFAYGISHHP
DFGDIISSQQEINRL
E
RPCNVSG*RI\SIEEI
LGPVEERLALHVL
HVESRPLYSPLQPD
RKSRWIRVIPFPRFF
DVILDDLSLTGEK
WMIGFEEHKQKTD
EGNFNWRFIFPFDY
AKKDAFWRLDKT
FQIWDNDKFSFDD
FLHQSQKTVVVKN
TLIFYEIEIFGEPAT
VELYDHDTYGADE
SLERMPRLAWFPL
LLASFELIQREKPA
DTDLPYPPPQREA
KPALQRTAIEILA
SYQLANISSPSLVVE
CVIRNLRKNPNFDI
LPREELYCPPITVK
RPVVGQCTIRSLES
SPSPQGGPDDVSL
DIDDKEPLIPIQNYA
SCYSRSQRSVSRCP
PRPGRPSGRSGLLA
GCFQCPPCLQRGL
APAHKGLFSGSLW
RAAATS*DRRAG
PSGTPRGEHLS*EN
AHQTAPQESQTS
-
EEISOUREII SSAMLI

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
6303	36671	A	6353		8655	EAAKDLADIAAFFRSGFRKNDE MKAMDVLPILKEKVAYLSGGR DKRGGPILTFPARSNHDRIRQE DLRRLISYLACIPSEEVCKRGFT VIVDMRGSKWDSIKPLLKILQE SFPCCIHVALIIKPDNFWQKQRT NFGSSKFEFETNMVSLEGLTKV VDPSQLTPEFDGCLEYNHEEWI EIRVAFEDYISNATHMLSRLEEL QDILAKKELPQDLEGARNMIEE HSQLKKKVIKAPIEDLDLEGQK LLQRIQSSESF
6304	36672	Α	6354	1	1338	
6305	36673	С	6355	152	404	
6306	36674	A	6356	137	1278	MDSQPNSTRGQSGRR/MEIKGI QLGKEEVKLSLFADDMIVYLEN A\TSQPKISLS**ATSAKSQDTKS MCKNHKHSYTPITDKQRAKS* VNSHSQLLQRE*NT*ESNL*GT* RTSSRKTTNHCSMK*KRIQTNG RTFHAHG*EESIS*KWPYCPR*F IDSMPSPSSYQ*LSSQNWKKRL SSSYGTKKEPASPSQS*AKRTKL EAS/LLPDFKLYYKATVTKTAW DMDEAGNHHSQQTITRTKNQT PHVLTHRWELNNENTWTQEGE HHTPGLVVRRSPDPEQNFKAVR CLDLPD/PSSSFLAVLTFHFKPA FLQSLR*DKPRS/SPALGTWHKP *KITPKKQ/HQAITKPNTEYRTA VKPSVVGALPDINSFLKLGGLH RITNGCLLD

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
6307	36675	A	6357	1	2569	MRTKTQHTRISWDAFKAVCRG
0507	150075	i `	0337	1	2309	KFIALNAHKRKQERSKIDTLTS
				-		QLKELEKQEQTHSKASRRQEIT
				1		KIRAELKEIETQKSLQKINESRS
		1				WFFERINNIDRPLARLIKKKREK
1		1				NQIDTIKNDKGDITTDPTEIQTTI
		1				REYYKHLYANKLKNLEEMDKF
		ŀ				LDTYTLPRLNQEEVESLNRPITG
	l	1				SEIVTIINSLPTKKSPGPDGFTAE
		Ī				FYQRYKEELVPFLLKLFQSIEKE
						GILPNSFYEASIILIPKPGRDTTK
	1					KENFRPISLMNIDAKILNKILAN
				}		l e
			ļ			RIQQHIKKLIHHDQVGFIPGMQ
						GWFNIRKSINVIQHVNRAKDKN
	1					HMIISIDAEKDFDKIQQPFMLKT
						LNKLGIDGTYLKIIRAIYDKPTA
						NIILNGQKLEAFPLKTGTRQGCP
1						LSPLLFNIVVEVLASAIRQEKEI KAQNLLKLISNFSKVSGYKINV
						l ~
			1			QKSQAFLYTNNRQTESQIMSEL
	}					PFIIASKRIKYLGIQLTRDVKDLF KENYKPLIKEIKEDTNKRKNIPC
1						·
						SRVGRINIVKMAILPKVIYRFNA IPIKLPMTFFTELEKTTLKFIWN
						QKRAHIAKSILSQKNKAGGITLP
						DFKLYYKATVTKTAWYSYQNR
						DIDQWNRTEPSEITPHIYNYLIF
						DKPEKNKQWGKDSLFNKWGW
				ŀ		ENWLAIWRKLKLDPFLTPYTKI
						NSRWIKDLNVRPKTIKTLEENL
						GITIQDIGMGKDFMSKTP/TSNG
						NKRQN*QMGSNETKELLHSKR
6308	36676	A	6358	1702	2252	FLSLFFFISLASSLSILLIHIMNSW
0308	30070	A	0338	1702	2232	IPHSPIFTDLNVESSRLCPLGDIM
				ľ		M*IPLTWYLIVTCTHLSREITTV
						PRGLARLWSFTRICPRTHCKPIP
	İ					A\VLLRAGCPPSCWWDQDTHS
						QLPSKRSSYTLFFCTVTEYPFLD
}		1				*WP*NHGTPC*TAPLWSMF/PCS
						CWYPVTGPGVSCSPSCVQVFSL FISHL*VRTCGVWFFVLAIVC*E
						*WFPASSMSQAVLVTVAL*YSL
						KSGSVMPPA\FSFGLGLTWRCR
ŀ						LFFGSIRTLK*IFPIL*RKSLVA* WGWH*IYKLRWAVWPFSRY*F
		1				FLPMSM/VMFFHLFVSSFISLSS
						GL*FSLKRSFMSLVSWIPKYFIL
						FEAIVNGSSLMIWLSVCLLLLH
						KNACDFCTLILYPETFLKLLISL
		1				RRFWAETMGFSRYTIMSSANR
		<u>L</u>	L	<u> </u>	<u></u>	DNLTSSFPN

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
ĺ	sequence	ļ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
6309	36677	Α	6359	207	1209	IPRHIHPPKTKPGRS*/TP*IDQ*Q
						ALKLRQ/CINSLPTKKSPGPDGF
1						TAKFYQSPSLTGKNDKEPEILPY
				,		FHADFLVVPGTHQASPVLMVL
						GHKSFPHRNSPPSVCMANFFTO
						KPPSQSIHNAVNTHTDPPLFLVL
						LKDRGLLVIWCHESQGDELWH
						-
						LVLLDSKCCFGSMTHRMEWSG
						PSAAGLLEFAGGPLQTLFAWVS
	1					AVEAAEQRILLNSKCCCLIIPLE
						ALSQRGIWPWQGSPVPRPWTTT
						TLWPVRNQATQQETSLMPCQL
						EGKEPLALPTKEAEIEGRRTGEI
						KKVERDSNTNKEELLLDLAFK
		1				VSSLFDKLEEVVELLVIGELQL
		1				YPSK
6310	36678	Α	6360	544	678	AFFHVSFGCINVFF*EVSVHML
	:					CPLFDGVVFFLVNLFEFFVDSG
6311	36679	Α	6361	7	94	VRTSGVWFFVLAIVC*E*WFPA
İ				}		SSMSLIY
6312	36680	A	6362	381	431	ATKQDSVSKKKEEEEEEEEE
						EEEEEEEEEEEEEEKEK/GNK
						ERKK/DKGKARQVQARQIYLLS
						LHLGRIRKVPNNNKA*SSCP
6313	36681	В	6363	1	603	
6314	36682	Α	6364	1	987	
6315	36683	В	6365]1	444	
6316	36684	Α	6366	977	1177	RTSISSPVSSTCSPLVMPTLLPQP
						TVPWRMWRPCTQ*RNAGTFSP
				ļ		AVAMGLPVEAVICLSV*DSPA
6317	36685	Α	6367	1148	1395	
6318	36686	Α	6368	33	663	VHVLCGDTDETGNHHSQQTIA
						RTKNQTPHVLTHRERLSLRRGL
						HSCAYSLHRWLGARHAQKGSR
						NTKGCGGQGPRTSERGPASGEF
					ł	SENCTSLLGQRGVFSPGRRALL
						HSPSDG\SVPLGAVDSSP\PASTR
						RDWSHWAVGKPSPEREPAGKV
						T*TTRQLQPASP*REHPLLNSQH
1						GHCLKISLPLLPTVLSPSPPLSLC
						SNVTFSTRPLT
6319	36687	A	6369	1	1314	
6320	36688	Α	6370	120	410	
6321	36689	A	6371	1	492	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
6322	36690	A	6372	3	945	EEVPRSMSKKKKRKRRKSRKR EEEEEEEEEEEEEEEEEEE EERR\EGEGEEEEEEEEEE
6323	36691	В	6373	1	792	
6324	36692	С	6374	348	884	
6325	36693	Α	6375	1	588	
6326	36694	В	6376	1	5339	
6327	36695	В	6377	520	3146	
6328	36696	Α	6378	1	2044	
6329	36697	Α	6379	1	585	
6330	36698	Α	6380	1	2433	
6331	36699	В	6381	53	441	
6332	36700	Α	6382	2	425	
6333	36701	A	6383	1386	2569	FLRKHRESGKDEEMTALCSFPG VAGEDELQVIQPEKSVSVAAGE SATLRCAMTSLIPVGPIMWFRG AGAGRELIYNQKEGHFPRVTTV SELTKRNNLDFSISISNITPADAG TYYCVKFRKGSPDDVEFKSGA GTELSVRAKPSAPVVSGPAVRA TPEHTVSFTCESHGFSPRDITLK WFKNGNELSDFQTNVDPAGDS VSYSIHSTARVVLTRGDVHSQV ICEIAHITLQGDPLRGTANLSEAI RVPPTLEVTSQPI\RAENQANVT WQVSNFYPRGLQLTWLENGNV SRTETASTLIENKDGTYNWMS WLLVNTCAHRDDVVLTCQVEH DGQQAVSKSYALEISAHQKEH GSDITHEPALAPTAPLLVALLLG PKLLLVVGVSAIYICWKQKA

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,	
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide	
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)	
6334	36702	A	6384	1	2453	MEPAGPAPGRLGPVLCLLLAAS	
						CAWSASRQLPAVQVLRGDVTC	
					1	GSDHVFLLAYIKHLAVAGRRA	
						LKATVHVHQRTLLRPGGCGSV	
		-				PGKRAVTLISKCSDGQPAFPGA	
						PAGNTNTFLVCSCGFPDLQSSR	
						GPERALCAFIQCLMVRPEEQRE	
						KNWPGPRGRSMARPGWNAKPP	
						VVQAKAFKNWAVLELSTVSAR	
						NLTGSRIPRENTKVILIGTIWVL	
						CFLGSALGVCDPGGTHSHLAM	
						HVVIFGPVGSNNPSIVIIIIIIRPIL	
						QIKSQRSRVGETFPGVSWVAEL	
						ESKSKHTFSQCQALLQPWGRK	
						LSSCSQRTYNLLLVCIQSMNVI	
						DKHLRKHRGSRVAGEEELQVI	
						QPDKSVLVAAGETATLRCTATS	
							LIPVGPIQWFRGAGPGRELIYNQ
						KEGHFPRVTTVSDLTKRNNMD	
						FSIRIGNITPADAGTYYCVKFRK	
						GSPDDVEFKSGAGTELSVRAKP	
						SAPVVSGPAARATPQHTVSFTC	
						ESHGFSPRDITLKWFKNGNELS	
			}			DFQTNVDPVG\ESVSYSIH\STA	
						KVVL\TREDVNSQSFW*GPTVT	
			}		;	LQGDPLRGTANLSETIRVPPTLE	
						VIQQPVRAENQVNVTCQVRKF	
						YPQRLQLTWLENGNVSRTETA	
						STVTENKDGTYNWMSWLLVN	
						VSAHRDDVKLTCQVEHDGQPA	
						VSKSHDLKVSAHPKEQGSNTA	
			}			AENTGSNERNIYIVVGVVCTLL	
6335	36703	A	6385	1	405		
6336	36704	A	6386	91	2926	PACPSPRLPFTNTDTHIDTSSPA	
0330	30701	1	0500	ļ''	2,20	SRSTRCLCISTHGVLAQSGGSSG	
						GPAVPTVQRGIIKMVLSGCAIIV	
						RGQPRGGPPPERQINLSNIRAGN	
						LARRAAATQPDAKDTPDEPWA	
						FPAREFLRKKLIGKEVCFTIENK	
						TPQGREYGMIYLGKDTNGENIA	
						ESLVAEGLATRREGMRANNPE	
						QNRLSECEEQAKAAKKGMWSE	
						GNGSHTIRDLKYTIENPRHFVD	
						SHHQKPVNAIIEHVRDGSVVRA LLLPDYYLVTV	
6337	36705	A	6387	2	448	DEFELDITEALA	
6338	36706	A	6388	1	645		
6339	36707	A	6389	1	294		
UJJ7	١٥١٥٢	1^	80209	11	1274		

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
6340	36708	I A	6390	23	321	RGREHTQGIPGAGKRWH*FKV
0340	30700	``	0550			LLTQ/CW*SEFSVIRQFKAGPGG
						LDQFRDRSK/RWLRDLRGRLLG
						SLNASLVLRALFSGNWGLTAN
						QL*LAVGAKGDPEA
6341	36709	Α	6391	1	1599	
6342	36710	Α	6392	1	561	
6343	36711	A	6393	1	1833	STLASSEGQGGFWRGPRALILL
						LLFLGPCHARSHDGKYSREKN
						QPKPSPKRESGEEFRMEKLNQL
						WEKAQRQERYPCRRRRRVKQV
						AAGTPRVFLKTPEMGVFELCRP
						SGPHYLISNAGVRLDLLNSPVL
					-	SIPESLRGKVNDLQNKKELLER
						KGRGETGLARPMHDLTAAQGG
						YGCALVHGAQVEGAGAQGWA
						GTVHATCRGETGTACQEALVT
						ATACSSCQRGQTRAPQGQHTL
						ESMGLNCLPQGAASGSDHHCG
İ						PRISDLPGSAARTSRWPASSELH
						LPPVRLAELHADLKIQERDELA
						WKKLKLDGLDEDGEKEARLIR
						NLNGHWRAEDAHQDGGVNK
						WDIPTVVGVGVAAIALVAVKL
ĺ						FFLPINIRCVLSESPATLHSLQV
						ASVLWSLEFLPYGSRSPVVIQET
						VILAKYGLDGKKDARQVTSNS
						LSGTQEDGLDDPRLEKLWHKA
İ						KTSGKFSGEELDKLWREFLHH
						KEKVHEYNVLLETLSRTEVHPG
						PQDGVASTGREHSPMETLFQEE
						LKHFEAKIEKHNHYQK\QLEIA
						HEKLRHAESVGDRRSV*SASPR
						ESTPCLEGRTKGAGLPR*KKHL
(2::	26512	ļ	620.4		105	QD\LSGRI\SRARHKRTLKALGS
6344	36712	Α	6394	1	185	DFFDNSAQVTIAGIPCDIRHVSP
						RKIECTTRAPGK\V*GSPPLSQAI
6245	26712	 	6205	,	1512	EGFFLKLEMLLRDWN
6345	36713	A	6395	1	1512	OCLUENTIN AND CALLED AND
6346	36714	Α	6396	117	268	QSLLLEYLHLYLHLCYLHLANK
						QELTLTWYKE*CTFFLPSV*NPF
6247	26715	_	6207	1	500	HQNLA
6347	36715	Α	6397	1	500	

	SEQ ID NO:		j .			Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide	of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	sequence		07/340,217	sequence	or peptide sequence	deterion, (-possible nucleonae insertion)
				,		
6348	36716	Α	6398	1301	2565	TLLTILPFYLFKNRTSLTGYLRI
						WILSTKLFCLSTLWLAPTLFLIP/
						LPDPHDCISLIHLAFTPFPHTSFF
		İ				PIPHPDHTWFIDGSSTWPNRHSP
						AKVGYAIVVSSTSILEATALPPS
						TTSQQAELIALTRALALAKGLH
						VNIYTDSKYAFHILHHNAVIWA
						ERGFLSTQRSSIIIASLIKTLLKA
						ALLPKEAGVIHCKGHQKASDPI
		ļ				AQGNAYADKTKFLDLAFPPLR
						LSFTCQITQAVSQALGIQWNLH
						TPYRPQSSGKDWTVFLPLALHR
						IRARPREATGYSPFELLYGRTFL
						LSPNLVPDTSPLGDYLPVLWQA
						RQEIHQAANLLLSTPDSQLYED
						TLAGRSVLIKNLTPQTLQPRWT
		ĺ				GPYLVIYSTPTAVRLQDPPHWV
						HRSRIKLCPSDNQPNLSSSSWKS
						QVLPTTSLKVTLISEEQ
6349	36717	A	6399	581	2517	LFLGYSHTSLLPFPPVQSLLHLL
						PLYLHTLTHKYGIPLLPPWRPIM
						HPLPSH*NLITLTLLNANIPSCST
						L*KD*SLLSLTCYSMAF*/QPINS
	:					PYNSPILSVQKPDKPYSSFRYVY
						SLFVESPTITIVPGPDFNLASHIIP
				•		DTTPDPYDCISLIHLTFTPFPHVS
						FFPVPHPFHTWFIDGSSTRPNCH
						TPAKAGYAIVSSTSIIEVTTLPHS
						TTSQQAELIALTRTLTLAKGLC
						VNIYTDPISCTTMLFYGLKCFLT
						MQGSSIINASLIKTLLKAALLPK
						EAGVIHCKGPQKASDPIAQDNT
						YADKVAKKAASVPTSIPHSSFS
						PSHLVTPTYSPTETSTYQSLPTQ
				•		GKWFLDQGKYLLPASQAHSILS
						SFCNLFHVGYKPLARLSELLISF
1						PSWKSILKEITSQCSICYSATPQ
						GLFRPPPFPKHQAWGFVPTQD
						WQIDFTHMPQVGKLKYLLVW
						VDTFTGWVEAFPTGSEKATVVI
						SSLLSDVIPRFCLPTSIQSNNGLA
						MISQITQAVSQALGIQWKLCTP
						YHPQSSGKVKWANGLLKTHLT
						KLSLQLKKDWTVLLPLALLRIR
						ACLRNATGYSPFELLYRCSFLL
						GPSLIPDTRPTWTAPPKTWHPY
						YLLSSHTPIHRSQLLIHALLLFTL
						PVYTVSPSHHS

OSSIBLE NUCLEOTIDE ENUCLEOTIDE ENUCLEOTIDE ENUCLEOTIDE ENUCLED EN CONTROLLE EN CONT
QVIAEGNICKIQE GEKDPESLLLPPL TYYNISVAKAELL AEIGLGEEEVDH ESISRKLSVLREA NSALGEEVEANL EKYHLFVGDLDK RLARVENALNSI /LIEKKQQLTGQL VDRREKLVFGMV DYQHFVKMKSA IKLGEEQLKCLR
GEKDPESLLLPPL TYYNISVAKAELL AEIGLGEEEVDH ESISRKLSVLREA NSALGEEVEANL EKYHLFVGDLDK RLARVENALNSI //LIEKKQQLTGQL VDRREKLVFGMV DYQHFVKMKSA IKLGEEQLKCLR
GEKDPESLLLPPL TYYNISVAKAELL AEIGLGEEEVDH ESISRKLSVLREA NSALGEEVEANL EKYHLFVGDLDK RLARVENALNSI //LIEKKQQLTGQL VDRREKLVFGMV DYQHFVKMKSA IKLGEEQLKCLR
GEKDPESLLLPPL TYYNISVAKAELL AEIGLGEEEVDH ESISRKLSVLREA NSALGEEVEANL EKYHLFVGDLDK RLARVENALNSI //LIEKKQQLTGQL VDRREKLVFGMV DYQHFVKMKSA IKLGEEQLKCLR
YYNISVAKAELL AEIGLGEEEVDH ESISRKLSVLREA NSALGEEVEANL EKYHLFVGDLDK RLARVENALNSI VLIEKKQQLTGQL VDRREKLVFGMV DYQHFVKMKSA IKLGEEQLKCLR
AEIGLGEEEVDH ESISRKLSVLREA NSALGEEVEANL EKYHLFVGDLDK RLARVENALNSI /LIEKKQQLTGQL VDRREKLVFGMV DYQHFVKMKSA IKLGEEQLKCLR
ESISRKLSVLREA NSALGEEVEANL EKYHLFVGDLDK RLARVENALNSI /LIEKKQQLTGQL VDRREKLVFGMV DYQHFVKMKSA IKLGEEQLKCLR
NSALGEEVEANL EKYHLFVGDLDK RLARVENALNSI /LIEKKQQLTGQL VDRREKLVFGMV PDYQHFVKMKSA IKLGEEQLKCLR
RLARVENALNSI //LIEKKQQLTGQL //DRREKLVFGMV PDYQHFVKMKSA IKLGEEQLKCLR
/LIEKKQQLTGQL /DRREKLVFGMV /DYQHFVKMKSA IKLGEEQLKCLR
VDRREKLVFGMV DYQHFVKMKSA IKLGEEQLKCLR
VDRREKLVFGMV DYQHFVKMKSA IKLGEEQLKCLR
DYQHFVKMKSA IKLGEEQLKCLR
IKLGEEQLKCLR
*I*VHSSPTQRLS
SLSLPAGLVGGD
RPLCSWLDPGKL
GGRSGAGTLQEG
PPLKPAIQTQSG
SWVIDWD
HIPHLGVLL/SAH
*MGSRTEGSATG
GPYDPADPMVLE
PHDPLAKFLFPVP
LVPEGGTLPPGD
KIAT*TLWVPPTF
IWKLPRLGASTL
FSHDWSGWDAG
GYNTCGCFHRL
MVQAVNGSTTL
INTACMANDOLLE

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
6369	36737	A	6419	38	530	HSPRRSAASLLSQRDHEPHQKE
0001						ETPNTSEHQKEKTPDTPPLRTV
						TLTVRVHGFILEVSETKNPPIPD
						TKLRG/ERLQGFGLAGFRSE/GC
						RPSP*VLQLIKAVWTQRPLGGR
						WDWAPWSRRWCSSGRLRPHR
						NQWSGWEAQAWGLQVPEPCP
						AGRQLRPGEKSSTAP
6370	36738	Α	6420	1	649	
6371	36739	Α	6421	1	987	
6372	36740	Α	6422	1	924	
6373	36741	A	6423	1	435	
6374 6375	36742	A	6424	1	147	
6376	36743 36744	A	6425 6426	3	913	
6377	36745	A	6427	1	159	
6378	36746	A	6428	313	1083	
6379	36747	A	6429	1	1245	MAQELRDTCTSFSSRFDQVEER
0317	30717		10125	`		VMVIEDQINEMKQEEKFREKR
						VKRNKQSLQEIWDYVKRPNLR
						LIGVPEKNLEETDKFLDTYTLPR
						LNQEEVESLNRPITDSEIEAIINS
						LPTKKSSGPDGFTAEFYQRYKE
						ELP/RQRHNKKREF*TNIPDEHR
						CKSPQ*NTGKPNPAAHQKAYPP
						*SSGLHPWDARLVQHTQINKRN
						PAYKQNQGQKPHDYLSRCRKG
						L*QNSAALHAKNSQ*IRY*WDI
						SQNNKSYL*QTHSQYHTEWAK
			İ			AGSIFFENWHKRGMPSLTTPIQ
						HSVGSSGQGNQAGERNKGYSI
						RKRGSQIVPVYR*HNCISRKPH
						HLSPKSP*AGKQ/SSAKSQD/DKI
						NVQKSQAFLHTNNRQTESQIMS
						E/APIHNCYKENKISRNPTYKGC
						EGPLQGELQTTAQRSKKGHKQI
6380	36748	Α	6430	39	314	KSEČLIYTKĠSQRLASNGTKLD
						GE*L**VDRIRLQKIGNDKLLPA
						KGGCSNPSQRS*KPRKKIRQMA
						N*NKQHRGDLK*PDGAENHGT
		<u> </u>				RTT
6381	36749	Α	6431	213	843	VDRSRLQKVSNNKLL*AKGECS
						NPLQGS*KP*KKIRRMAN*NKH
						CREDLK*PDGVENNGTRTS*FM
			1			HKLQ*PI*SSGRKGISD*R*N**N
						KIHQG*NEGKSVKGSQRERSSY
				-		PQRKAHQTNSGSLSRNPTSQKK
						VGANIQHS*RKEFSTQNFISSQT
						KLHK*RRNKIFYRQADAERFCH
						HQACVTRAPEGSTKHGKKQPV
			<u> </u>	<u> </u>	<u> </u>	PAPAKTGQIV

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
		<u> </u>				
6382	36750	Α	6432	1	2112	MRAFVVLMDVSKLPSERGGPG
						LRSNQQHMRMFLTSVDRGSVS
						LLADAVAAATPQGTKSQPRGPS
						NLAADLRVAEESSGMENLFLK
						MSFDSIRPAQDGQCTPEISMVTP
						SPGWSVHTGDLDGHPLPRMVS
						AHRRSRWSPPPQDGQCTPEISM
						VIPSPGCSVHTGDLDGHPLPRM
						LSAHRRSRWSPPPQDAQCTPEIS
		1				MVTPSPGWSVHTGDLDGHPLP
						RMVSAHRRSRWSPPPQDAQCT
		<u> </u>				PEISMVTPSPGCSVHTG
6383	36751	Α	6433	1	648	
6384	36752	Α	6434	1	1539	MLVSFVSLGSLCLQPGSQTLLE
						KNRTVKPHVSFTLLPALSHVSE
		1				KNEAESMNSLIPPPPNLHTPAQ
						APFPLPTKEQDRSSSPATEQSW
						TENDFDELTEVGFRRSVITNSSK
	1	ļ				LKEDVRTHCKEAKNLEKRLHE/
	1					SAN*NQQCREDLK*PEAEIHGT
		1				RTS*HMHKLQ*PI*SSGRKGVSD
						*GSNQ*NK*WGEWNQVGKHSA
			,			GYYPGELPQPSKAGQHSNSGNT
						ENTTKILLEKSNPKTHNCQIHQ
						G*NEGKSVKGSQRERTREARA
						NKFKS*QKARYN*DQCRTERD
						RNTENPSKN**IQEPVF*KDQQN
						**NASKTNKEEKREESNRRNKK
				1		**KNRTVKPHVSFTLLPALSHVS
		ŀ				EKNEAESMNSLIPPPPNLHTPAQ
						APFPLPTKEQDRSSSPATEQSW
		•				TENDFDELTEVGFRRSVITNSSK
						LKEDVRTHCKEAKNLEKRLHE
	-					WLTRINSVEKTLNDLKLKSMA
			1			RELHDTCTSFNSRFDQVEERVS
						AIEDQTNEINNGENGTKLENTL
		1				QDIIQENFPNLARQANIQIQEIRR
		Ì			1	TPQRYSSRKATPRHIIVRFTKVE
		1				MKEKVLRAAREKGLERQEQTN
	1					SKASRRQDITKISAELKEIETQK
						TLQKINEFRSRFFKKINKIDKTL
						ARLIKKKREKNQTDAIKNDKR
						DITTDPTEIQTTIREYYKHLYAN
	1	1				KLENLEEMDKFLDTYTLPRLNQ
	1					EEAESLNRTITGSEIEAIINSLPT
6385	36753	В	6435	137	586	
6386	36754	Α	6436	129	506	
6387	36755	Α	6437	1	237	
6388	36756	Α	6438	182	1034	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
6389	36757	A	6439	203	650	SAIVCGSPLRLRMQAVSSRINSA
						HSRGAIYSQASSSAACPSVSIAF
			İ			LRR*IEANVS/WIRTLLSTLCTPK
						CGPGFKHRIVLCKSSDLSKTFPA
						AQCPEESKPPVRIRCSLGRCPPP
	-	1				RWVTGDWGQ/CDSRDPRPGRR
						RRDFCRNGRAGAGH
6390	36758	Α	6440	207	426	AECTRCAIPK/HSCYPVLLIVVS
						P*KKQ\NCSLGELSCWRRESCRS
						SWSNKALPSLTRCLRGFVCGLS
						CYKGSS
6391	36759	Α	6441	524	783	TAVLAARISNQWILSCWS/AVG
		}				CTPKCGPGFKHRIVLCKSSDLS
					}	KTFPAAQCPEESKPPVRIRCSLG
						RCPPPRWVTGDWGQLSPTREK
6392	36760	A	6442	1	1524	
6393	36761	A	6443	1	8738	
6394	36762	A	6444	1	642	
6395 6396	36763 36764	A	6445	1	549 434	
6397	36765	A	6446 6447	8	2922	
6398	36766	A	6448	18	1158	FLDFDCPYGSAHITGLRFWLPW
0378	30700		0440		1130	YRKRQRYWGSHGNCMPPPSLR
						VPERCPSHTEPRNLTGA*ELLLL
		ŀ				GLSEDPELQPILAGLSLSMYLVT
						VLRNLLISLAVSSDSHLHTPMC
						FFLSNLCWADIGFTSAMVPKMI
:						VDMQSHSRVISYAGCLTQMSLF
						AIFGGMEE\NMLLSVMAYDWF
		1				VAICHPLHYPVIMNPHLGVFLD
						LGAFFLSLLDSQLHSWIVLQFTF
						FKNVEISNFV*DPSQLLNLACSD
						SVINSIFIYLDSIMFGFLPISGILW
		1				SYANNVPSILRISSSDRKSKAFS
						TCRSHLAVVCLFYGTGIGVYLT
						SAVAPPLRNGVVASVTYAVVT
						PMLNPFIYSQRNRDIQSALRRLR
						SRTVESHDLLHPFSCVGKKGQA
6399	36767	Α	6449	2	1535	
6400	36768	A	6450	1	635	
6401	36769	A	6451	222	1040	
6402	36770	Α	6452	2	1135	
6403	36771	Α	6453	1	576	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
6404	36772	Α	6454	2	694	QCGGIQRSHGSWSQEASDSVW
	1					QLQSMWMLDKLTGVFAPRPST
	1					GPHKLR\ECLSRIIF\LRNRLKYA
						LTGDEVKKI\CMQRFIKI\DGKV
1			ļ			RT\DITYPVG\FMDVI\SIDKTGR
						EFPVLIYDTKG/RAFAVHRITP\E
			[EAQVPSCAKVRRDLCGAQKGIP
		ļ				HLVTH\DARTIPLPRSPSSKVND\
						TIQI\DLEDWARFT*FSSKFDNW
	1					*TCCMGDWEGA*P*GRNWVLI
						HQQRRRAPWDLF
6405	36773	Α	6455	1	243	
6406	36774	Α	6456	3	766	
6407	36775	Α	6457	1	741	
6408	36776	В	6458	500	8643	
6409	36777	Α	6459	1	7863	MKPRTLTVRVTALKVARLESV
						PSDVQMCSEFLPSDSAAQLASP
						SGSRTGAAGGAAYQSCAVLPH
						SSALWVVDGTRCRGAGGSAHQ
		ı				GGSATQEPTERVEGSGMAGCR
						SRVLPHWKAANVLGMSVESAP
				,		AVEEEKGEELEQKEKEKEEDTS
}						GNTTHSLGAEDTASSQLGFGVL
		1				ELSQSQDVEENTVPYEVDKEQL
						QSVTTNSGYTRLSDVDANTAIK
						HEEQSNEDIPIAEQSSKDIPVTA
					:	QPSKDVHVVKEQNPPPA
6410	36778	Α	6460	2	669	ERRRKLTIPCPPPHPAPPTGGQ
					:	ACRENAEGIL*AANTSEPGKDA
						EK*KIKV*LAP/CSHSGSSLQSDP
						HFGCSLGPS/DPALGLSGCILPPC
						SGSYLAPAPNSRIPKPGLKPW/H
						/ESYLPSAVRPHGTQAPGLWPE
						RGL*PAGFLLRPEPVRAASER/Q
						EAVSPER*VPRPPKAQQLHGQS
						PDPSSLCTALRQDHTHWLDARS
						VYPDWRGQPWPPLHQDCGHG
					:	GWR
6411	36779	В	6461	241	369	
6412	36780	A	6462	1	510	
6413	36781	В	6463	1	630	OCD CVODULL BUVCCUVCCO
6414	36782	Α	6464	229	561	SCRGYSRILLLFHKSSWCSQVIS
		1				IFRYNIICSRLPRSWKWRLPVY
						MNWL*YHWFKQYHSSINFISFR
						SMWFIIINFKRSPAFILSGSWSIS
L						IARWQIFVFFTTFGSRTKV

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	•	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
6415	36783	A	6465	1271	[1710	L TEGLGHRSSETPGGAALDALTG
						ERPRVACFPAARGRGLRPRPRL
						MIGPFETGGGQRLLTLRPRRRR
		ŀ				QRRRP/WQGLERPHVRNWGIEL
		ŀ				ETLRDAFRPQPSVWRTARGEA
				:		GKRKGRLVECLGSQRKVLEGV
				•		LLYAVTRGSFPNLFTPRD
6416	36784	В	6466	1	772	
6417	36785	Α	6467	1	1272	
6418	36786	Α	6468	3	236	
6419	36787	С	6469	1	457	
6420	36788	В	6470	148	324	
6421	36789	A	6471	1	1407	
6422	36790	Α	6472	1	651	
6423	36791	Α	6473	619	835	
6424	36792	Α	6474	52	535	
6425 36793	Α	6475	319	647	LCRRIYMYFCVMKNWMQSLW	
						KLCTNKR*RIDKPTLHQVVVRK
						*LTPLGMKNQVDSLELLETRSL
					•	KKDPVELGSKSLRRCCCQTEPV
						NLLPHVSIGTLKI*SRILRLQRP
6426	36794	Α_	6476	39	515	
6427	36795	В	6477	1	807	
6428	36796	Α	6478	3	247	GANICCW*GLQE/VFHSW*KTQ
						EPSWLHLVDPTTGLQVELPASP
						APFARIPQPLGGRWDWAPWSR
(120	0.5505	 	6.450	0.7	2.62	GWCSSGRLGQHRSPWSG
6429	36797	A	6479	27	363	
6430	36798	В	6480	1132	1268	
6431	36799	A	6481	1628	2008	
6432 6433	36800 36801	A A	6482 6483	953	1087 443	QLAGRCGGRGMSGNRGC\WGA
0433	30801	A	0483	-	443	CGPAGVPGG/CWAWCALHSEE
		1				PASPALRSISRRALAAFPRGRAR
						DLQPAMPEPPTPSVGSCAAQAS
						PTSAAPCSMAPSPIDHPRAEECV
		}				ARRGTGRHDELIVALPFMMIPW
						HSSIQPFTIHPFNWNFG
6434	36802	С	6484	175	423	110010111111111111111111111111111111111
6435	36803	В	6485	122	608	
6436	36804	A	6486	644	1149	KTMHWKAPLMSWHMHRASSQ
0430	30001	ľ.	0.00			KQTRTGTQT/MEAQLQQLQRY
						QEALLQVLRKGEKKAFNMGNI
						SEVLQRADESCSQFYERLYEAF
						QLCSPFDPEATENQCMVNAVF
						VGQTQGDIRRKLQKLECFTGM
						NVTQVLEVATKVYTNCEEETQ
						GKGQFAGGSTYGKGGECPKTK
6437	36805	Α	6487	1	2847	CATE OF A CATE OF THE CATE OF
6438	36806	В	6488	329	742	
0	1		1	1	1	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion) .
6440	36808	A	6490	13	630	INPPPPPFRPEPPSSSNSPKMTDH KGERGEATRYMFSRPF\RK\HG VVPLGPHICRI\YRKGDIVD\IKG
-					, 	MGYCSKKGIAPTKCY\HGQNLE GVLPMLPQHACWPFVVNQPVL GQSFFPRELMCRIEHI*ALLRAR
						DKLPETVLKENDSEKERSPNEK GTWGSNLKRHPGSTPEEAHFV
						RTQWGRSLELAGTLFPMNFMG IIGVKEIK
6441	36809	Α	6491	1	1020	
6442	36810	Α	6492	1	330	
6443	36811	С	6493	250	372	
6444	36812	Α	6494	1	345	
6445	36813	A	6495	2	335	TVRAYNVPQHAVGIVVNRQVK GKIPAKRINVGIKHFKHSTS*EN FLKRMKENDQKKKEAKEKVT WIQLKHQSAPPRE/AQCERTNG KEPELLEPLPYEFMHNTYIHIYV CVH
6446	36814	Α	6496	2	490	
6447	36815	A	6497	150	536	NCKISFLHFCYIFVKALKRISAL SRGKILAKRINVRIEHIKHSKSR DSFLKRVKENDQKKKEAKEKG TWVQ/RKAPGKNLVYISLVLRA L*G*DLTHHIIILFPFFFL**PAPP REAHFVRTNGKEPELL
6448	36816	Α	6498	54	553	
6449	36817	Α	6499	131	803	TMFCQTGSSVRRVGPGPWAEN RTAMLRPCGGTFARKTRLSRLG NSPKMNEQQRGKRRGTPIIMFS KGLLRKHG\VVPLAHIFMR\IYK\KGDIVRHPRGMGYLVPKRELP PQSCYPWPKPGRSLTNVTPAML VGHLL*NKQV*GQRFLPKRNL MWRI*APFRHFLRSPRLASLKT CGRENGSRKKERKPKGGKVTW GFQLKRRHLGFPPQEEATFLLK EPIGGREP
6450	36818	Α	6500	1	489	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
6451	36819	A	6501	29	2028	ENEEEGEKMASTDYSTYSQAA AQQGYSA\YTAQPTQGYAQTT QQAYGQQSYGTYGQPTDVSYT QAQTTATYGQTAYATSYGQPP TGYTTPTAPQAY\SQPVQG\YGT GAYDTTTATVT\TTQASYAAQS AYGTQPAYPAYGQQPAATCPT RTED*\NKPTETSQPQSSTGGYN QPSLGYGQSNYSYPQVPGSYP MQPVTAPPSYPPTSYSSTQPTSY DQSSYSQQNTYGQPSSYGQQSS YGQQSSYGQQPPTSYPPQTGSY SQAPSQYSQQSSSYGQQSSFRQ DHPSSMGVYGQESGGFSGPGE NRSMSGPDNRGRGRGGFDRGG MSRGGRGGGRGGMGSAGERG GFNKPGGPMDEGPDLDLGPPV DPDEDSDNSAIYVQGLNDSVTL DDLADFFKQCGVVKMNKRTG QPMIHIYLDKETGKPKGDATVS YEDPPTAKAAVEWFDGKDFQG SKLKVSLARKKPPMNSMRGGL PPREGRGMPPPLRGGPGGPGGP GGPMGRMGGRGGDRGGFPPRG PRGSRGNPSGGNVQHRAGDW QCPNPGCGNQNFAWRTECNQC KAPKPEGFLPPPFPPPGGDRGRG
		Hey Y				GPGGMRGGRGGLMDRGGPGG MFRGGRGGDRGGFRGGRGMD RGGFGGGRRGGPGGPPGPLME QMGGRRGGRGGPGKMDKGEH
6452	36820	Α	6502	1	693	
6453	36821	A	6503	3	657	
6454	36822	A	6504	1	434	-
6455	36823	A	6505	208	443	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
6456	36824	A	6506	3	2650	LLPAFLLLCSDSHFLQFCSQFAP RKTQGSGSVNNIQANIIAKKSN NHLMKKNDQKFKDLPQMAGH PIHHQAQGPPGAKNPQEQQRRP MGQTTLPAEQEESRVKCKNCG AFGHSARNKTCPIKRWSGALPL QVLGSHKEKENLKPAKAQLPFT TPGPFTTNDREKERSPSSTMNPS EMQRKASPRRWKHHNQTPSIH KVNKMVMSEEQMKLPSTKKA EPPTWAQLKKLTQLPKKSLEN\ TKVTQTPENTLLAALMIVST/A GAAAANYTYWAYVPFPPLIRA VTWMDNPIEVYVNNSAWVPGP TDDRCPDKPEEEGMMINISTGY RYPPICLGRAPACLMPAIQNWL VEVPTVGTTSRFTYHTGLLTFR DVAIEFSQEEWKCLDPALEDSY TRRKANSCGKPYKCNECGKAF TQNSNLTSHRRIHSGEKPYKCS ECGKTFTVRSNLTIHQVIHTGE KPYKCHECGKVFRHNSYLATH RRIHTGEKPYKCNECGKAFRGH SNLTTHQLIHTGEKPFKCNECG KLFTQNSHLISHWRIHTGEKPY KCNECGKAFSVRSSLAIHQTIHT GEKPYKCNECGKVFRVGRGVSI GTTCLLSVFQVITVSSRKSRASP ENRATQSILILAPPTSRGSGPGP ASVLRRLAQTRKRMAWTESCT AACAFPSCLVLLYRGRGVPTDL QVPAPLFSTQSKLKRPGSRTKSS
6457	36825	A	6507	54	360	CSTMNPSEMQRIAPPRRQRHRS RAPSAHKMNRMVMSEEQMKL PSTKKAEPPTWAQLKKLTQLA KKK\LENTKVTQTPENMLLAAL KTVSTVSAGVPSSSEESD
6458	36826	A	6508	525	3656	SSTMNPSEMQRKAPPRRRRHR NRAPLTHKMNKMVTSEEQMK LPSTKKAEPPTWAQLKKLTQLA TKYLEN\TKVTQTPENTLLAAL MIVSMVVSLPMPAGAAAANYT YWAYVPFPPLIRAVTWMDNPIE VYVNNSAWVPGPTDDHCP\AK PEEEGMMINISTGYRYPPICLGR APACLMPAIQNWLVEVPTVGT TSRFTYHTIDLMTEKWVVLASV EVLLRFPLKPGEDPTARYVSNK KCQPSVDWPTTISQRRGY
6459	36827	Α	6509	1	246	
6460	36828	Λ	6510	1	246	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
6461	36829	Α	6511	1	279	
6462	36830	Α	6512	1	363	
6463	36831	С	6513	109	245	
6464	36832	Α	6514	2	429	LMTFCWDTPRQSGVPRERMPH
						TGTWRTVGIRCPKKKLRSANSS
						MMRGKVQPFRP/DIQAYGAAPF
						EDLQVDFREMPKCGGNKYVLV
		İ				LGRTYSGWVEAYPTRTEKTRE
						VTPVLLRDLIRRFRPPLWIGSDN
						GPAFLAALVQKTAK
6465	36833	С	6515	31	895	
6466	36834	Α	6516	2	4735	
6467	36835	Α	6517	263	418	
6468	36836	Α	6518	120	295	YDPLPTWILGKISQRGAEEPAQ
						HPGCQL*LPPP/SPPRTSSSPSCA
		<u> </u>				GLTSVSPRPRFPR
6469	36837	Α	6519	1	1758	MKIRRFPSSLQLSADLQVPVR
	3				KAPRQGVSQEKRTKPSLELMIS	
		į				GTISQEDIRNNVTRGCTPPAIQR
	1		ļ			VISSSPLLAIRNNVTEGVYTPCD
						TGGRVHLPVILLLISRDGGHDIS
		l				FNIAVDVHSPCDTDPNIQGVEY
						DMTPNIAMNVQPPGTISQKRCT
						LPAILGLISYASSVNIRSNITGWL
						YIDRYVGSHVILYPLGIRMGVT
						GEMKGERRRGLHAATKWLEE
						HAPADYQNPQEYGRTQLPGTQ
						PQLDPHEREDMQRLNRDREAL
						LEGFKRGAQKATNVNKVSEVI
		ł				QGKEESPAQFHQRLCEACAKQ
						VLVNGDAVSREEKRKENERQA
						RRNADLLVSCSNQSGPPKEARE
						ELWTKDYRPGQDLRLLSQATL TFHPTVPSPSTLLGLLPAEDSWF
						TCLDLKDAFFPIRSAPESQKLFA
						FQWEDPESALAKTVRQRCVSC
					:	RQHHA\GKVQPFRP/DIQAYGA
						AAFEDLQVDFTEMPECGGNKY
		1				LPVLGRTYSGWVETYPTRAEK
						AREVTRVLLRDLIPRLELPFRIG
]		SDNGPAFVADLLQKTATVLGIT
						RKLHAASRPQSSGKGIQNNRTG
						GVYTPCDIESHVILFRSGY
6470	36838	A	6520	1111	1342	MFVPHIGFQNTAALCLLRLGVL
04/0	30030	\ <u>`</u>	0.520	' ' ' '	1.374	FP*PVGFPAGSCLLI*WNRHWH
			1			VPDFSCVTTPQEWCGGVSDVC
						CGHPHAEPFHLQ
<u></u>		<u> </u>	<u> </u>	<u> </u>	1	COMMENTING

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	}	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
6471	36839	A	6521	1	1144	MRGAQKATNVNKLSEDIQGKE
		İ				ESPTQFYERLWEAYRMYTPFDP
		i				DSPENQRMIPMALVRQSAEDM
						RRKLQKQAGLAGMNPSQLLEI
						ASQVFVNRDAVSRKENGKENG
						GQARRYADLFSRTKNYQPVQD
					İ	LRLLHQAKLTFHPTVPNPSTLL
						GFPPAEDSWFTCLDLKDAFFPIR
				}		LAPERQKLFAFQWEDPESGWPP
ĺ						CWRALAATALLVQEANKLTLG
ļ						QKLNIKASRAVVTLMNTKGHH
						WLTNATLTDYQTLLCENPRITIE
						VCNTLHPATLLPVSKSPVKPGC
						VEVLDSIDSSRPDLWDQPWASV
						DWELYLDGSSFFNPQGEVEG\S
	1					RGDTSELPPCWVCGIPALTQRL
						EKQHLPPSGHQGSLKHLIWDLL
					,	LLTKKRTFSSMI
6472	36840	A	6522	1	554	ELTRICH SSMI
6473	36841	В	6523	63	620	
6474	36842	Α	6524	1	796	SRTKNYQPVQDLRLLHQAKLT
				ł		FHPTVPNPSTLLGFPPAEDSWFT
						CLDLKDAFFPIRLAPERQKLFAF
						QWEDPESGWPPCWRALAATAL
				:		LVQEANKLTLGQKLNIKASRA
				1		VVTLMNTKGHHWLTNATLTD
						YQTLLCENPRITIEVCNTLHPAT
						LLPVSKSPVKPGCVEVLDSIDSS
			1			RPDLWDQPWASVDWELYLDG
						SSFFNPQGEVEG\SRGDTSELPP
1						CWVCGIPALTQRLEKQHLPPSG
						HQGSLKHLIWDLLLLTKKRTFS
6475	36843	A	6525	3	2640	
6476	36844	Α	6526	2745	3732	IRIGKNYFKVHMEPKKSPHRQV
1	Ì					NPKPKEQSWRHHTT*LQTILQG
						YSNQNSMPGPTPQ\PSTPAPGG
						NLRNPQSSDLLQVTKQQGQAL
		1				AIQREAPLHRIPAPEAIPWYFQP
ļ						QPATQLGSPPVDPPSSAMM/SR
						RAHRSRGPDRQGYPLQGD*PGE
						PRPQEH*RGHSQERLPSSEKQTP
						ICPPAQATQHPEEPDAHQPYKH
				!		LFQVCAHQGHPVAQRRN*PGY
						WERYHSAEDPELQPILAGLSLS
						MYLVTVLRNLLIS
6477	36845	A	6527	1	1860	
6478	36846	Α	6528	1	1458	
6479	36847	В	6529	1	1503	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence	ŀ	09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
6480	36848	Α	6530	3	515	TISQKRCTLPAILGLISYASSVNI
						RSNITGWLYIDRYVGSHVILYP
İ		ŀ				LGIRMGVTGEMKGERRRGLHA
		1				ATKWLEEHAPADYQNPQEYGR
						TQLPGTQPQLDPHEREDMQRL
						NRDREALLEGFKRGAQKATNV
ļ					 	NKVSEVIQGKEESPAQFHQRLC
			•			EAYRMYTPLDHDSPENQRMIH
		ŀ		:		MALVRQSAEDMRRKLQKQAG
				-		LAGMNPSQLLEIASQVFVNRDA
						VSPKENSKENGGQARRHDDLF
						SPVEPDCVEVLDSIHSSRPDLRD
						QPWPSGDW/RTICGWEQLLQPP
						RRERCRVCSDNPGH/WLLKPHR
			ŀ			CPRPLQARKLNSLLSLGP*NSTR
						DCVRPIVCILPLITIALKISA
6481	36849	В	6531	96	2524	
6482	36850	В	6532	61	2025	
6483	36851	Α	6533	674	2913	
6484	36852	В	6534	120	1849	
6485	36853	A	6535	1	426	
6486	36854	A	6536	2	475	
6487	36855	Α	6537	1	501	
6488	36856	Α	6538	15	593	SMWWNSAREPCPWRIVDDCGG
		1				AFTMGVIGGGVFQAIKGFRNAP
		1				VCRLLSEAPLFIYSCSRSVSPTV
		ł				NVSSERAESRPTLFMAVSLHMA
		l				WCLAHIGIRHRLRGSANAVRIR
		1				APQIGGSFAVWGGLFSTIDCGL
Ì		i				VRLRGKEDPWNSITSGA\LTGA
						VL\AARSAPLAM\VGSAMMGGI
		<u> </u>				LL/ALIEGVSILL/TR*TATV
6489	36857	Α	6539	2	557	RRFRASAMEEYAREPCPWRIVD
						DCGGAFTMGVIGGGVFQAIKG
		l				FRNAPVGIRHRLRGSANAVRIR
		1				APQIGGSFAVWGGLFSTIDCGL
						VRLR\GKEDPWNSIT\SGA\LTG
						AVL\AAR\SGPLAMGGLQQ*WG
						GILFAPHLRAFGILLTRYTAQ\Q
						FRNAPPFLEDP\SQ\LPPKDGTPA
		<u> </u>				\TGYP\SYQQYH
6490	36858	A	6540	223	2274	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide inscrtion)
6491	36859	A	6541	2795	3762	FLHFHLRNA VPHQQWNKAGW
		ŀ				RMTLRS*EKKASDDQITLSYGR
						TFKPK/DKEVENFEQNLEECITR
		l				ITNTEKCLKELMELKTKARELR
		Ì				EECRSLRSRCDQLEERISVMED
						EMNEMKREGKFREKRIKRNEQ
						SLQEIWDYVKRPNLRLIGVPES
						DGENGTKLENTLQHIIQENFPN
						LARQANVQIQEIQRTPQRYSSRI
						AIPRHIIVRFTKVEMKEKMLRA
1						AREKGRVTLKGKPIRLTADLSA
						ETLQARREWGPIFNILKENNFQ
						PRISYPAKLSFISEEEIKYLTDKQ
ŀ	İ					MVRDFVTTRPALKELLKEALN
ŀ						MERNNRYQLLQNHAKF
6492	36860	Α	6542	245	1293	TGAVPIRPSWQRWGKNRTEKL
					ETLKRRAPLLLQRNAVPHQQQ	
					1	NKAGWRMILTS*EKKASDDQIT
					ł	LSYGRTFKPK/DKEVENFEKNL
İ						EECITRITNTEKCLKELMELKTK
						APELREECRSLRSRCDQLEERV
ļ						SAMEDEMNEMKREGKFREKRI
		1				KRNEQSLQEIWDYVKRPNLHLI
						GVPESDVENGTKLENTLQDIIQ
l						ENFPNLARQANVQIQEIQRTPQ
						RYSSRRATPRHIIVRFTKVEMK
						EKMLRAAREKGRVTLKGKPIR
]						LTADLSAETLQARREWGPIFNIL
						KEKNFQPRISYPAKLSFISEGEIK
						YFIDKQMLRDFVTTRPALKELL
						KEALNMERNNRYQPLQNHAK
6493	36861	С	6543	116	283	
6494	36862	Α	6544	51	281	KSQNQLLSSKGEYADGQQDQV
					1	SIPAPSPYTLSGPQTQSCPVRPE
						LGL*GASSAVQAPAVRYLTRFL
1						LFVMSNLHSP
6495	36863	В	6545	1	897	
6496	36864	В	6546	91	8030	
6497	36865	A	6547	126	362	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
6498	36866	IA	6548	936	2064	ERADCLFKRKTNNQKGHLHRK
	1					PICTSPSSKTKSR*NHKDGEKTE
						QKNWKL*NAERLSSS/IRNAVP
						HQQQNKAGWRMILTS*EKKAS
		İ				DDQITLSYGRTFKPK/DKEVENF
						EKNLEECITRITNTEKCLKELME
						LKTKARELHEECRSLRSRCDQL
						EERVSAMEDEMNEMKREGKFR
		j	;			EKRIKRNEQSLQEIWDYVKRPN
		l				LRLIGVPESDVENGTKLENTLQ
						DIIQENFPNLARQANVQIQEIQR
						TPQRYSSRRATPRHIIVRFTKVE
						MKEKMLRAAREKGQVTLKGK
						PIRLTADLSAETLQARREWGPIF
						NILKEKNFQPKISYPAKLSFISEG
						EIKYFIDKQMLRDFVTTRPALK
						ELLKEELNMERNNRYQPLQNH
6499	36867	A	6549	1	767	MAGAPPPASLPPCSLILDCCASN
0499	30007	^	0349	1	/0/	QRDSVGVGPSEPGVGYSLVVR
						RFLSRSEKRNIRVGVTRFSSEM
						NPVPQMEMQKSPSSASLTLGA
		1				VDRSCSYSAILAPPLGFCFYPLY
ŀ						ENSTLQSAKKRDAELANGALGI
						IELNNDYTLKKVMKPLITSNTV
Ì	:					TDEIERANVFKMNGKWYLFTD
						SRGSKMTIDGINSNDIYMLGY/D
						IKLFNRPLQAAEQNRACAANGS
						*SKRCDIHLLSLRSAASQRQQC
						GYHKLHDKQRLLRG
6500	36868	В	6550	50	327	G I HKLHDKQKLLKG
6501	36869	A	6551	333	1301	FLHFHLRNAVPHQQWNKAGW
0301	30809	^	0551	333	1301	RMTLRS*EKKASDDQITLSYGR
						TFKPK/DKEVENFEQNLEECITR
						ITNTEKCLKELMELKTKARELR
						EECRSLRSRCDQLEERISVMED
						EMNEMKREGKFREKRIKRNEQ
				:		SLQEIWDYVKRPNLRLIGVPES
						DGENGTKLENTLQHIIQENFPN
						LARQANVQIQEIQRTPQRYSSRI
			1			AIPRHIIVRFTKVEMKEKMLRA AREKGRVTLKGKPIRLTADLSA
						ETLQARREWGPIFNILKENNFQ
						PRISYPAKLSFISEEEIKYLTDKQ
						MVRDFVTTRPALKELLKEALN
(500	2.070	 	(550	252	2204	MERNNRYQLLQNHAKF
6502	36870	A	6552	352	2304	
6503	36871	Α	6553	1	1617	<u> </u>

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
6504	36872	A	6554		257	MGPIHKISHYVYANISKSEKVL
050-1	30072	``	0334	[]		KSEHFRSQAFQPGQQSETVSEK
	ļ.					KKKTRKKRKKRKKKKKK
						KKKKKKKKKKKKKKKKK
						SE*DRLGKKEEDEEEEEEEEE
		1				EEEEEEEEEEEEEEEE
						EEEEEEEEEEEEEEE
6505	36873	A	6555	5	156	RKKGGG/EREEGEGEEEEEE
0303	30073	`	0333			EEEEEEEEEEEEEEVVL
						LTPGTGAEDFKKH
6506	36874	Α	6556	li	279	ETT GTGALDTAKKT
6507	36875	A	6557	3	120	TEEEEEEEEEEEEEEE
0307	30073	`	0557		120	EEEEEEEEEEEEEEEEEE
6508	36876	A	6558	3	123	CECCEDECE CECED CECED
6509	36877	A	6559	239	432	CLWLFQEEEEEEEEEED*EEE
0309	30077		0337	237	432	EEE/EEEEEEEEEEEEEEEE
		ŀ				EEEEEEEKIFLGHRVGI
6510	36878	A	6560	1	588	ELECEPEERII EGIIKVGI
6511	36879	A	6561	l i	204	
6512	36880	A	6562	75	234	
6513	36881	В	6563	100	2510	
6514	36882	A	6564	47	4976	
6515	36883	A	6565	3	4755	SCRCRRRRRGPAPAMAQILPVR
0313	30003		0303	[7733	FQEHFQLQNLGINPANIGFSTLT
						MESDKFICIREKVGEQAQVTIID
		l				MSDPMAPIRRPISAESAIMNPAS
						KVIALKAGKTLQIFNIEMKSKM
						KAHTMAEEVIFWKWVSVNTV
		1				ALVTETAVYHWSMEGDSQPM
						KMFDRHTSLVGCQVIHYRTDE
						YQKWLLLVGISAQQNRVVGA
						MQLYSVDRKVSQPIEGHAAAF
						AEFKMEGNAKPATLFCFAVRN
						PTGGKLHIIEVGQPAAGNQ
6516	36884	В	6566	47	482	1 TOOKETHIE VOOT AAGITQ
6517	36885	A	6567	1	555	MNARPHKVD/GRVVEPKRAV*
0517	30003	\^_	0307	1	333	EDSQRP\AFVAGIKEDTEEHHLR
						DYFQQYGKMEVIEIMTDRGSG
						KKRGFAFVTFDNHDSVDKTVV
]		QKYHTANDHNCEGRKALSKQE
						MASASSSQRGHSSSGNFRGGHG
			1			1 - 1
		1				GGFGGNDIFGYGENFSGHGGFS
						GSCGGGGYGSSGGGYNEFGND
6510	26006	 _	(500	1	C45	GSNTGGGGSYNDFGN
6518	36886	Α	6568	1	645	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
6519	36887	Α	6569	2	944	GRGLRKLFIGGLSF*FIDESLKS
03.7	30007	, , , , , , , , , , , , , , , , , , ,	0507	-		QVGEWGTLTDCVVMRDPNTK
						RSRGFGFVTYAT\VEEVDAAMN
						ARPHKVDGRVVEPKRAVSRED
						SQRPGAHI/TLVKKIFVGGIKED
						TEEHHLRDYFEQYGKIEVIEIMT
						DRGSGKKRGFA\FVTFDDHDSV
						DKIVIQKYHTVNGHNCEVRKA
						`
						LSKQEMASASSSQRGRSGSGNF
						GGGRGGGFGGNDNFGRGGNFS
						GRGGFGGSRGGGGGGGGGGG
						NGFGNDGSNFGGGGSYNDFGN
						YNNQASNFGPMKGGNFGGRSS
						GPYG\GGGQYF\AKPR\NQGGY
		<u> </u>				GGSSSSSSYGSGRRF
6520	36888	A	6570	1	1512	
6521	36889	A	6571	2864	3677	RCQLDHLPCHLCCCCHR/CIPSL
						RDPQQAPGSTRLSRAPHIESRV
						GRKPPEEPANPTMNSLTLRDKQ
			i			/HDASTC/DWKRAL*VPESGLPR
						ALLKYIRCPNMSSAQKPRELSK
						RRN*K*PCVKRTPFRPT*E/DLP
						YSGKKTGVKGLICPVPGTAVK
						APQRPPGPQQPHPAPQSASETA
						WFPVADRRVFSENPGWTAAKT
						LGTLSSPCQAGAGVGRQDGDP
						GQELTSQEKPWVGRKIPEHAV
						ADKGYRQQCGAGEARGGPVF
		<u> </u>				WAAAPRTSWQQASYT
6522	36890	Α	6572	1	666	
6523	36891	A	6573	2926	3304	EYSLGHRHYNWKC*ALGQKDQ
						VTFATKFAPTSHRTRKQPELWR
	-					/QDPGSYGEKQPAWCSVPPGRF
						SGGLVGLGFLDGGIVVQPLRRP
						CCQDCVCGFPTLRSSTKTSTEL
						GPSFLPCERQGKETISVT
6524	36892	Α	6574	1	665	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
6525	36893	Ā	6575	3	1121	SSTFPRMPWRLVTSWTAYIPEL
						EGSTSDPDSPAVAQKGEKPVKL
İ		l				GNSLWGSHTEPVPRPGTGPRPL
						SFVHRLPTMDPHKVNALRAFV
						KMCKQDLSVLHTKEMHFLRE
						WVESMGGKLSPATQKVKSEEN
						TKEEKPDSKKVEEDLKADEPSS
						EESNLLIDNEDVIEPDPDAPQEM
						GDENAEITEEMMDQANHKKVA
		1				AIEALNDGELQKAIDLFTDAIKL
						NPQLAILYAKRASVFIKLQKPN
						AAIRDCDRAVEIY\SAQPYKWR/
						GKYEQKREEREIKERIERVKKA
		Ė				REEQERAQREEEARGQSVAQY
						GSFPDGLPGGMLEMRGGMPGK
		:				PGITGLNEIFSDPEVLAAVQDPK
						VMV/AFFQDVAQNPANMSQYQ
						GNPKVM
6526	36894	A	6576	3	1351	PEAGGERERERAQREEDSTRQS
0320	3007-	``	0570	[1331	RAQYGSFTGGFPGGMPGNFPG
						GMPGMGGPSARSRRAQRLSLE
}						PDRSPPSYAHHLPTMDPRRVNE
						LRAFEKMCKQDPSVLHTEEMR
						FL\RECVESMCGKVPPATHKAM
						SEENTTEEKPDSEKAEEDLQAD\
						EPSSEESDLRKLIKEGVIEPDTD
			ŀ			APQEMGDENAEIPEE\MMDQA
		1				NDKKVAAIEALNDGELQKAIDL
						FTDAIKLNPRLAILYAKRASVF
						VKLQKPN\AAIRDCDRAIEINPD
						SAQPYKWRGKAHRLLGHWEE
			1			AAHDLALACKLDYDEDASAML
						KEVQPRAQKIAEHRRKYERKR
1						EEREIKERIERVKKAREEHERA
						QREEEARRQSGAQYGSFPGGFP
						GGMPGNFPGGMPGMGGGMPG
						MAGMPGLNEILSDPEVLAAMQ
						DPEVMVAFQDVAQNPANMSK
						YQSNPKVMNLISKLSAKFGGQ
6527	36895	A	6577	2	433	GPPLNLSSPRGGILKTYGCELCG
3321	30073	<u>۱</u>	5577	_	133	KRFLDSLRLRMHLLAHSAGAK
						AFVCDQCGAQFSKEDALETHR
						QTHTGTDMAVFCLLCGKRFQA
						QS\ALQQHMEVHAGVRSYICSE
						CNRTFPSHTALKRHLRSHTGDH
			-			PYECEFCGSCFRDE
6529	36896	_	6578	1	687	FIECETCOSCFRDE
6528 6529	36896	A A	6579	2	449	
0329	الجووردا	/A	03/3	<u> </u>	לדיד <u>ן</u>	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
:	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
6530	36898	IA	6580	852	1084	TDAGCYYAARQCGTAVPSRDV
	20070	'				HIRHQGTPVPQF*TAPLRARS/E
						VSPARRRARMLSSSASFASANIT
						TSSAWSDTLQP
6531	36899	A	6581	1	681	
6532	36900	Α	6582	1	483	
6533	36901	A	6583	1	273	TRGPWCDSVLRGCSLEQRSFIS
						VRLLSYLSACRHPMEDSMDMD
					}	MSPLRPQNYLFG\SLGAGAKDE
				1		LHIVEAEAMNYEGSPIKVTLAT LKMS
6534	36902	A	6584	1	678	DKINO
6535	36903	A	6585	3	634	
6536	36904	A	6586	4	479	CSAKTAIRGVSECLIHHHGIPHS
						TASDQGTHFTAKEVWQWAYA
						HGIHWSYHVLHHPEEA/GKER
						WNGLLKSQLQCQLGDNTLQG*
						GKVLQQAVYVLN*HPIYGTVSP
F						IARIH/GSRNQG/VETELTPLTITP
						SDPLAKFLLPVPPTLHSAGLEIL
						VPEGEIR
6537	36905	Α	6587	1	1539	MVGKAKWKPLELPLPRKMVSQ
						KQYRILGGTAEISVTIKDLKDA
						RVVIPSTSPFYSPIWPVQKIDGS
						WRMTEDYRKLNQVVTPTAAA
						VPDVVSLLEQINTSPGTRYAAID
						LTNAFFSIPFHKAHQKLFAFSW
	,					QGQQYTFTVLPWGDINSPALCH
						NLIRRDPDCFSLPQDITLVHHID
				ļ.		DIMLTGSREQEVADTRDFLVKD
<u> </u>						KLLHLAPPTTKKQAQHLVGLF
						GFWRQHIPLLGVLLQPIYRVTQ
		1			}	KVASFEWSQEQKTALQQVQAA
						VQAVLPLGPYNPADPMVLEVS
						VADRDAVCSLWQAPIGESQWR
						PLGFWSKALPSSADNYSPFERQ
						LLA/CHKVGHAQQHSISKWK*Y
						IHDRAQADYHPWT\QACLIHCQ
						GIPHSIASDKDAHFTTKEVWQW
						THAHEIHWCYHVLHHPEAAGL
						TERWNRLLKSQLQRQLGDNTL
						QGSGKVLQKAMYALNQHPIYG
						TISPIARIHGSSNQGVEVAPLTIT
						PSDPLAKFLLCVPTTLRSAGLD
			<u> </u>			VLVPEGGMTQQQFH

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
6538	36906	A	6588	1	566	MGRNQSRKAENSKNWSASSPP KDHSSLPAMEQSWMENDFDEF TEVGFRRSVITNFSELKEHVLTY CKEAKNVEKRIYRSRNQGMEV KVAPLTITPSDPLANFFASVPAT LHSAGLEVLVPEGGTLPLGDTT MIPLNWKLRLRTGYFGLLLRLS QQAKKGAMVLA/GVVDPDYQ NEISLLLHN*GKTHDH
6539	36907	A	6589	221	357	
6540	36908	C	6590	228	342	
6541	36909	A	6591	745	849	
6542	36910	A	6592	1	240	
6543	36911	A	6593	2	495	FVRLVGRGDCDPLLSVCLTTMP LYEGLGSGG\EKTAVVIDLGEA FTKCGFAGETGPRCIIPSGIKRA GMPKPVRVVQYNINTEELYSYL KEFIHILYFRHL\LVNPRDRRV\V IIES\VLCPS\HFREDTHSCFFFKY FE\VPSVLLA\PS\HLMALLTLGI NSWHGPRL
6544	36912	Α	6594	1	579	
6545	36913	Α	6595	1	621	
6546	36914	Α	6596	1	351	
6547	36915	Α	6597	1	447	
6548	36916	Α	6598	114	880	
6549	36917	Α	6599	1	2625	
6550	36918	Α	6600	1	1674	
6551	36919	A	6601	1035	2235	QTHSQYHTEWAKTGSIPFENW HKTGIPSLTTPIQHSVGSSGQGN QAGEGNKGYSIRKRGSQIVPVC RRHD/VYI*KTPLSQPEISLS**A TSAKS/LGYKINVQKSQAFLYT NNRQTESQIMSELPFTIASKRIK YLGIQLTRDVKDLFKENYKPLL KEIKEDTNKWKNIPCSWVGRIN IVKMAILPKVIYRFNAIPIKLPM PFFTELEKTTLKFIWNQKRARIT KSILSQKNKAGGITLPDFKLYY KATVTKTAWYWYQNRDIDQW NRTEPSEITPHIYNYLIFDKPEK NKQWGKDSLFNKWCWENWLA ICRKLKLDPFLTPYTKMNSRWI KDLNVRPKTIKTLEENLGITIQD IGMGKDFMSKTPKAMATKAKI DKWDLIKLKSFCTAKETTIRVN RQPTK

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
6552	36920	A	6602	369	1917	AGEGNKGYSIRKRGSQIVPVCR RHD/VYI*KTPLSQPKISLS**AT SAKS/LGYKINVQKSQTFLYTN NRQTESQIMSELPFTIASKKIKY LGIQHTRDVKDLFKENYKPLLK EIKEDTNKWKNIPCSWVGRINI VKMAILPKVIYRFNAIPIKLPMT FFTELEKTTSKFIWNQKRARITK SILSQKNKAGGITLPDFKLYYK ATVTKTAWYLYQNRDIDQWN RTEPSEMTPHTYNYLIFDKPEK NKQWGKDSLFNKWCWENWLA ICRKLKLDPFLTPYTKINSRWIK DLNVRPKTIKTLEENLGITIQDI GMGKDFMSKTPKAMATKDKID KWDLIKLKSFCTAKETTIRVNR QPKKWEKIFATYSSDKGLISRIY NELKQIYKKKTNNPIKKWAKD MNRHFSKEDIYAAKKHMKKCS SSLAIREMQIKTTMRYHLTPVR MAIIQKSGNNRCWRGCGEIGTL LHCWWDCKLVQHILTHRWEL NNEITWTQEGEYHTLGTVVGW GEGGGIALGDIPNAR
6553	36921	Α	6603	1	2781	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
6554	36922	Α	6604	1320	2403	LQTILQVYSNQNSMVLVPKQR
				,		YRSMEQNRALRNNATYLQLSD
						L*QT*EKQAMGKGFPI**MVLG
			1			KLASHM*KAETGSLPYTLYKN
						QFKMD*RLKR*T*NSKKQSPSP
						PPKERSSSPATEQSWTENDFDE
				·		LREEGFRRSNYPELREDIQTKG
						KEEIQTTIREYYKHLYANKLEN
	-					LEEMDKFLNTYTLPRLNQEEVE
						SLNRPITGSEIVAVINSLPTKKSP
						GPDGFTAEFHQRYKVELVPFLL
			1			KLFQSIEKEGILPNSFYEASIILIP
						KPGRDTIKKENFRPISLTNIDAKI
						LNKILANRIQQHIKKLIHYDQV
						GFIPGMQDWFNIRKSINVIQHIN
						RTKVKNHMIISIDAEKAFDKIQ
						QRFMLKTLNKLAQNLLKLISNF
						SKVSGYKINVQKSQAFLYTNNR
						QTESQIMSELPFTIASKRIKYLGI
						QLTRDVKDLFKENYKPLLKEIK
						EDTNKWKNIPCSWVGRLNIVK
						MAILPKVIYRFNAIPIKLPMTFF
						TELEKTTVKFIWNQKRACITKSI
						LSQKNKAGGITLPDFKLYYKST
						VTKTAWYWYQNRDIDQWNRT
						EPSEIMPHIYNYLIFDKPEKNKQ
						WGKDSLFNKWCWENWLAICR
ļ						KLKLDPFLTPYTKINSRWIKDL
İ				5		NVRPKTIKTLEENLGITIQDIGV
ļ						DKDFMTKTPKAMATKPKIDKW
						DLIKLKSFCTAKETTIRAFDELD
İ						DVHPHEEIEACRVWQNYGTHP
6555	36923	Α	6605	2202	2515	KSLQCEHSSHIQDLPLACLGLR
						KHYMVAQYL*HSHS/ASGGGSR
						YDVLKISVSCSSTIDI*LAFVSLG
						*DKRTS**TNLAESSPNFFHWV
ŀ						WIFLASKKFCSVPAR
6556	36924	Α	6606	1934	2077	AIPFRIQAWARTSCPKHKKQWQ
						QKPKLTNGI*FN*RASAQQKKL
						PSE
6557	36925	В	6607	1	2325	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
6558	36926	A	6608	1524	2877	IRKRGSQIVPVCRRHD/VFV*KT
						PSSQPKISLS**ATSAKS/LGYKI
						NVQKSQAFLYTNNRQTESQIMS
						ELPFTIASKRIKYLGIQLTRDVK
						DLFKENYKPLLKEIKEDTNKW
						KNIPCSWVGRINIVKMAILPKVI
						YRFNAIPIKLPMTFFTELEKTTL
	1					KFIWNQKRARIAKSILSQKNKA
		:				GGITLPDFKLYYKATVTKTAW
						YWYQNRDIDQCNRTEPSEIMPH
						IYNYLIFDKPERNKQWGKDSLF
						NKWCWENWLAICRKLKLDPFL
						TPYTKINSRWIKDLNVRPKTIKT
						LEENLGITIQDIGVGKDFMSKTP
						KAMATKDKIDKWDLIKLKSFC
		ĺ				TAKETTIRVNRQPTTWEKIFAT
						YSSDKGLISRIYNELKQIYKKKT
						NNPIKKWAKDMNRHFSKEDIY
			ŀ			AAKKHMKKCSSSLAIRERQIKT
						TMRYHLTPVRMAIIKKSGNNR
						DMDEIGNHHSQ
6559	36927	Α	6609	1	2226	
6560	36928	Α	6610	1	674	MGKKQNRKTGNSKTQSASPPP
		1				KERSSSPATEQSWMENDFDELR
						EEGFRRSNYSELREDIQTKGKE
						VQNFEKNLEECITRITNTEKCLK
						ELMELKTKARELREECRSLRSR
						CDQLEERVSAMEDEMNEMKRE
						GKFREKRIKRNEQSLQEIWDYV KRPNLRLIGVPESDVENGTKLE
						NTLQDIIQEN/CPQSSKAGQRSD
						SGNTENATKILLEKSNSKTHNC
		ŀ				QIHQS
6561	36929	В	6611	1	5175	Q111Q3
6562	36930	A	6612	1666	2032	LQTILQGYSNQNSMVLVPKQR
0502	30750	1	00.12		2032	YRSMEQNRALRNNATYLQLSD
						L*QT*EKQAMGKGFPI**MVLG
						KLASHM*KAETGSLPYTLYKN
			İ			QFKMD*RLKH*T*NHKNPRRKP
		ļ]			RHYHSGHRHGQGLHV
6563	36931	Α	6613	1596	2169	AGEGNKGYSIRKRGSQIVPVCR
						*HD/VYI*KTPLSQPKISLS**ATS
						AKS/LGYKINVQKSQAFLYTNN
						RQTESQIMSELPFTIASKRIKYL
						GIQLTRDVKDLFKENYKPLLKE
						IKEDTNKWKNIPCSWVGRINIM
						KMAILPKVIYRFNAIPIKLPMTF
						FTELEKTKVHMEPKKCPHHQG
						NPKPKEQSWRHHTT
		<u> </u>				-

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
6564	36932	A	6614	1667	3103	AGEGNKGYSIRKRGSQIVPVCR *HD/VYI*KTPLSQPKISLS**ATS AKS/LGYKINVQKSQAFLYTNN RQTERQIMSELPFTIASKRIKYL GIQLTRDVKDLFKENYKPLLQE IKEDTSKWKNVPCSWVGRINIV KMAILPKVIYRFNAIPIKLPMPF FTELEKTILKFIWDQKRARIAKS ILRQKNKAGGITLPDFKLYYKA TVTKTAWYWYQNRDIDQCNR TEPSEITPHIYNYLIFDKPEKNK QWGKDSLFNKWCWENWLAIC RKLKLDPFLTPYTKINSRWIKD LNIRPKTIKTLEENLGITIQDIGM GKDFMSKTPKAMATNAKIDK WDLIKLKSFRTAKETTIRVNRQ PTKWQKIFATYSSDKGLMSRIY NELKQIYKKKTNNPIKKWAED MNRHFSKEDIYAAKKHMKKCS SSLAIREMQIKTTMRYHLTPVR MAIIKKSGNNRLTLLNNHNNPQ
6565	36933	A	6615	3134	3288	PLISIKEPREKEELNLQV
6566	36934	A	6616	1	2630	
6567	36935	В	6617	877	7936	
6568	36936	Ā	6618	2	1589	
6569	36937	A	6619	1	240	
6570	36938	A	6620	200	486	
6571	36939	A	6621	1	558	
6572	36940	Α	6622	1	3885	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
:	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
6573	36941	Α	6623	1	2302	KLPVTNRAAHGLCASLGPDFM
			•			TFKVPSSWRHSDSICREWFTLA
						GAPSRDVLLVSAIITVSLSVTVV
						LCGLCHWCQRKLGKRYKNSLE
	:					TVGTPDSGRGRSEKKAIKVSRP
						CPPPAKASATHPTCKSPRSQEK
						VESPGDLDRDFWNNNESTVQQ
	ļ					KWSSYPPKEFILNISPYAPYGDP
						RLSLNGTLLSGAKVAAAAGLA
						VEREGRLGEKPAPVPPPGEDAL
			:			RSGGAAPSEPGSGGKAGRGRW
						RTVQSHLAAGKLNLSNFEDSTL
						STATTLESIPSSTGEPKCQRPRT
	ļ					LMRQQSLQQPLSQHQRGRQPS
:						QPTTSQSLGQLQAHMASAPGP
						NPRAYGRGQARQGTSAGSKYR
:						AAGGRSRSNPGSWDHVVGQIR
						NRGLDMKSFLTTSSRAAKWNF
:						TNSETVPALDLDSAVVLIPWISS
						KEDTGDSHYGRVAIYHKGLSFS
						SDGLVYPGAKGREGRMVVLSL
:				:		VLGLSEQDDFANIPDLQNPGTQ
		ŀ				1
						QNQNAQGDKRLPAGGKAVNT APVPGQTPHDESDRRTEPRSSV
E						1
r	<u> </u>					SDLVNSLTSEMLMEDWPPAVF
						SSGWAIATAHPASLHSPLIGYSF
	1					LAARGGLLLFEADFCWCKRSG
:	ľ					STFPDCDGVANLAATEMALSQ
						LRFSKETAMRESLVMLSPGSEE
:						DEAHEGCSRENLGRIQFSVGYN
:						FQESTLTLKIMKAQELPAKDFS
(574	26042		((2))	1	57(GTSDPFVKIYLLPDKKHKLKTK
6574 6575	36942 36943	A A	6624 6625	1	576 155	
6576	36944	A	6626	4746	5258	DACAODSI DDVI IVHODSDEEDI
03/6	30944	'A	0020	140	0رےر	DAGAQPSLPRYLIYHQPSRFFRI TSTVTPFSKPISSLLWLV*/GLHG
						1
						DILL/RLLQS*RC*CRSHPEYHSG
						PPGRWQGPSPAPQKCSRTTSRS
						RSSR*HPPPPRPGGSAPSSPASYS
						WTPPRTVPGSWARRWRRHLSR
						PSWATQPSWTATLWSQPLWRL
655=	2604-	<u> </u>	6607		700	KWEDCLSPGRSRLQ
6577	36945	A_	6627	1	588	

SEQ ID NO:	SEQ ID NO: of peptide sequence	1	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
6578	36946	A	6628	444	1056	GPLLREPHQLCHFPVAGFVLDA GLPEAAAAPHDRGPELAGPQA AGQHPAQVAHQPRAAAQQRQ GRRAREEAGAQAQEEPRGPQE RRRLAEGLRRQCGHPAGRDG RGGYRRAAQGPARGGRTPEWQ QSGGARRTPSVGHPRRARAD\W CLTGANDALSCSSPTCGKFLKT VSKVPVCRRGLAVNRKEAEGR SSESPTSSATFQSPALSWMLVS QRLLLPHMIGAQSSPGPKRPGN TLRKWLTSPVRRLSSGKADGH VKKLAHKHKKSREVRKSADAG SQKDSDDSAATPQDETVEEDTV AQPRGRREGAGRQSGSRAAGP GGRRASAIRGGHARTDVGKRN GTRTRKELMPLHGKRKDAAEH DEPTSASRPAGERPELTLAAVQ TEYNCSFPS
6579	36947	Α	6629	128	1103	
6580	36948	Α	6630	1	602	
6581	36949	Α	6631	3	1618	
6582	36950	Α	6632	1	602	
6583	36951	Α	6633	3	1618	
6584	36952	Α	6634	1	912	
6585	36953	Α	6635	195	396	VLPLHRQI**LLH*SKKHNHYK GKQQDLR/CSSTVPRRPQGILPII SFSNVPVGSRRLKAPLVFIGPG

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SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
			<u> </u>	sequence		
6586	36954	Α	6636	2	1813	FNTPLSTLDRSSRQKVNKDIQD
						VISALHEAGLIDIYRTLHPKSTE
						YTFFSAPHHTYYKIDHIVGSKA
						LLSKCKRTEITTNCLSDDRAIKL
						ELRIKKLTQNYTTTWKLNNLLL
ł						SDYWVNNEMKAEIKMFFETNE
						NKDTMYQNLWDTFKAVCRGK
						FIAVNAYKRKQERSKIHTLTSQ
						LKEPEKQEQTHSKASRRQEITKI
						RAELKEIDTQKTLQKISVSRSW
l	İ					FSERINKIDRLIARLMKKKREK
						NQIDAIKNDKGDITIDPTEMQTT
						IREYYKNLYANKLENLEEMDK
						FLDTYILPRLNQEEVELPNRPIT
				İ		GSENEAIINSLP/TKKSPGPDEFT
						AKFYQRYKG\LARAIRQEKERE
						DTQLGKEEVKLLLFAEDMIVYL
					•	ENPIVSAQNLLKLINNFSKVSGY
						KINVQKSQAFLHTNNRQTESQI
			}			MSECPLTIASKRIKYLGIQLTRD
					:	VKDLFKENYKPLLNKIKEDT\N
						KWKNIPRSWLGRNNIVKMTILP
						KVI/YKKTTLNFIWNQERARIAK
					İ	TILSKKNKAGGITLPDFKLYYK
						ATVTKTARYWTLTIKKGSVWPI
						SLKSEKRGHRTRPLWKLCHND
						VQGELLKTIQFLFGSPLGHSSLD
6587	36955	Α	6637	1	2988	MPPHSEQPHSPSTRQKRKVPLF
						VRQCPSFVIQAPKAKMDTFDGR
						KKQFTRFLFLFSMDQLGQGRFS
						MQTLSFALAVYRKDSPLVACQ
						VQALGNLEPSSVEAHVSSHGIAI
						DRKGWHVSLFSIQPGDCFPKAL
						VEDSPRDRARRQQSSKEEGWC
						RDRRRLKFHSKKGAAAIVVKS
						KKYESPSFGVCFESLLNPPRLTS
						RREKTISSSKRCRQCHAEETTV
						VFWAKESQTGEQTGRGAGQRR
						MGMIICKACSMLPAE
6588	36956	В	6638	362	1463	
6589	36957	Α	6639	1	1061	
6590	36958	Α	6640	1	1581	

SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
6591	36959	A	6641	46	862	AAFPAPRTLFFLSRPLRPTACAA FRPEFWRLNMKLPARVFFTLGS RLPCGLAPRRFFSYGTKILYQN TEALQSKFFSPLQKAMLPPNSF QGKVAFITGGGTGLGKGMTTL LSSLGAQCVIASRKMDVLKAT AEQISSQTGNKVHAIQCNVKYP DMVQNTVSELIKVQGHPSTVIT N*STERSSISFLLPTIYA*DWFRF CSTQVLLPKQVWKPMSKSLAA
6592	36960	A	6642	1	2902	EWGKIWNAIQC MVKGSLQQEELTILNIYAPNTE APRFIKQVLRDLQRDLDSHTLI MGDFNTPLSTLDRSTRQKINKD TQELNSALYQADLIDIYGTLHP KST\EYTFF\SAPHHTYSQNWTT IVGSKALLSEHKRTEIIT\NYLSD \HSA\IKLELRIKKLTQN\RSATW KLNNLLLNDYWVHNDMKAEI KMFFETK\ENK\DTTYQNLW\DT FKAVCRGKFIAK\NAHNRKQE\ RSKIDTLTSQLKELEKKEQTYL KASRRQEIT
6593	36961	Α	6643	1	5127	
6594	36962	В	6644	1	3570	
6595	36963	A	6645	1	3663	
6596	36964	В	6646	143	3122	
6597	36965	A	6647	1	3235	
6598	36966	Α	6648	1	3249	
6599	36967	Α	6649	1	3297	

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SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
6600	36968	A	6650	1	2563	MKAEIKMFFETNENKDTTNON
				}		LWDAFKAEEVESLNRPITGAEI
						GAIINSLPTKKSPGPDGFTAEFY
						QRYKEELVPFLLKLFQSIEKEEI
	i					LPNSFYEASIILIPKPGRDTTKKE
						NFRPISLMNIDAKILNKILANRI
						QQHIKKLIHHDQVGFFPGMQG
		l				WFNIRKSINVIQHINRAKDKNH
	ŀ					MIISIDAEKAFDKIQQPFMLKTL
						NKLGIDGTYFKIIRAIYDKPTAN
						IILNGQKLEAFPLKTGTRQGCPL
						SPLLFNILLEVLARAIRQEKEIK
						GIQLGKEEVKLSLFADDMIVYL
						ENPIVSAQNLLKLISNFSKVSGY
						KINVQKSQAFLYTSNRQTESQI
	•					MSELPFTIASKRIKYLGIQLTRD
						VKDLFKENYKPLLKEIKEDTNK
						WKNIPCSWVGRINIVKMAILPK
						VIYRFNAIPIKLPMTFFTELEKTT
						LKFIWNQKRAHITKSILSQKNK
						AGGITLPDFKLYYKATVTKTA
				1		WYCYQNRDIDQWNRTEPSEITP
						HTYNYLIFDKPEKNKQWGKDS
						LFNKWCWENWLAIWRKLKLD
ŀ						PFLTPYTKINSRWIKDLNVRPKT
						IKTLEENLGITIQDIGMGKDFMS
						RTPKAMATKAKIDKWDLIKLK
						SFCTAKETTIRVNRQPTTWEKIF
						ATYSSDKGLISRIYNELKQIYKK
l .	.					KTNNPIKKWEKDMNRHFSKED
						IYAAKKHMKKCSSSLAIREMQI
						KTTMRYHLTPVRMAIIKKSGNN
6601	36969	Α	6651	1	3402	-
6602	36970	Α	6652	1	3288	
6603	36971	A	6653	1	3168	
6604	36972	Α	6654	1	3516	
6605	36973	Α	6655	1	3693	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
6606	36974	Α	6656	1577	3354	TEPKTKTT*LSQ*MQKRPLTKF
<u> </u>						NNLSC*KLSIN/IVLEVLARAIRQ
						EKEIKGIQLGKEEVKLSLFADD
						MIVYLENPIVSAQNLLKLISNFS
				1		KVSGYKINVQKSQAFLYTNNR
						QTESQIMSELPFTIASKRIKYLGI
		1				QLTRDVKDLFKENYKPLLKEIK
						EDTNKWKNIPCSWVGRINIVK
}						MAILPKVIYRFNAIPIKLPMTFF
						TELGKTALKFIWNQKRARITKSI
						LSQKNKAGGITLPDFQLYCKAT
						VTKTAWFPSGDVGLEADFSPSH
		1				TLKTQFFSCLAEFAAASCFFQR
]				MNGFGMAMTTTYSTGAAESPL
						PSCSIDQGDDTKLHRARSPGRT
						FPAAAGIPAAAAPDGPPLSLLH
		1				KLWFPVELGGRALPRAEESHGE
		1				VAALGVMVVAQGGKNQGEEA
		1				RSTPWLRPTSHLPPCSSSSAWW
						TEQTDAHPLLFLLCLGIYLLNA
						LSNLSMVALVRSDGALRSPMY
						YFLGHLSLVDVCFTTVTVPRLL
						AGLLHPGQAISFQACFAEMYFF
		ľ				VALGITESYLPAAMSYDRATAA
		ļ				CRPLRYGALVTPWALRLAARY
						DRLASVVYAVITPTLNPFINSLR
						NKEVKGALKRGLRWRAAPQE
6607	36975	Α	6657	1	3514	MELKTKARELREECRSLRSRCD
						QLEERVSAMEDEMNEMKREG
						KFREKRIKRNEQSLQEIWDYVK
						RPNLRLIGVPESDVENGTKLEN
						TLQDIIQENFPNLARQANIQIQEI
						QRTPQRYSLRRATPRHIIVRFTK
						VEMKEKMLRAAREKDRSTRQK
	:		1			VNKDTQELNSALHQADLIDIYR
						TLHPKSTEYTFFSAPHHTYSKT
						DHIVGSKALLSKCKRTEIITNYL
						SDHSAIKLELRIKNLTKSRSTTW
	10.50=6	ļ	5550		0016	KLNNLLLNDYW
6608	36976	A	6658	3	3316	
6609 6610	36977	A	6659	1	4794 3570	
6610	36978 36979	A B	6660 6661	1	3384	
6612	36980	A	6662	1	3429	
6613	36980	A	6663	1	3780	
6614	36982	A	6664	1	3894	
0014	20902	ΙΔ.	10004	11	JUJ4	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
6615	36983	A	6665	1	3335	MVKGSIQQEELTILNIYAPNTG
		ŀ				APRFIKQVLSDLQRDLDSHTLI
	İ	İ	ļ			MGDFNTPLSTLDRSTRQKVNK
						DTQELNSALHQADLIDIYRTLH
		ŀ				PKSTEYTFFSAPHHTYSKIDHIV
		ļ				GSKALLSKCKRTEIITNYLSDHS
						AIKLELRIKNLTQSRSTTWKLN
1					:	NLLLNDYWVHNEMKAEIKMFF
}						ETNENKDTTYQNLWDAFKAVC
						RGKFIALNVYKRKQERSKIDTL
						TSQLKELEKQEQTHSKASRRQE
]			•			ITKIRAELKEIETQ
6616	36984	В	6666	1	3145	
6617	36985	Α	6667	1	4398	
6618	36986	Α	6668	523	3852	
6619	36987	Α	6669	1	3934	MGKKQNRKTGNSKTQSASPPP
						KERSSSPATEQSWMENDFDELR
		l				EEGFRRSNYSELWEDIQTKGKE
						VENFEKNLEECITRITNTEKCLK
ļ						ELMELKTKARELHEECRSLRSR
						CDQLEERVSAMEDEMNEMKRE
						GKFREKRIKRNEQSLQEIWDYV
						KRPNLRLIGVPESDVENGTKLE
						NTLQDIIQENFPNLARQANVQI
						QEIQRMPQRYSSRRATPRHIIVR
						FTKVEMKEKMLRAARQKAPH
						HTYSKIDHIVGSKAL
6620	36988	В	6670	1	5215	
6621	36989	Α	6671	1	7171	
6622	36990	Α	6672	3	493	
6623	36991	A	6673	1	729	
6624	36992	Α	6674	3	800	
6625	36993	Α	6675	1	327	WGGUW EEDDDDWWA DAARDAAG
6626	36994	Α	6676	2	462	KSSWLEEDDDPVVARVNRRMQ
						HITGLTVKTAELLQVANYGVG
						GQYEPHFDFSRRPFDSGLKTEG
			[NRLATFLNYMSDVEAGGAT\VF
						PDLGAAIWPKKGTAVFWYNLL
			-			RTGEGDYR\TRHAACPVLVGCK
						WVSNKW\FHES\GQEFLRPCGST
			<u> </u>		<u></u>	EVD

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence		in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
6627	36995	A	6677	3	717	RHRLRNRLKDVGGEAFYPLLE
						DDQSNLPHSNSNNELSAAATLP
						LIIREKDTEYQLNRIILFDRLLKA
	1					YPYKKNQIWREARVDIPPLIRG
						LTWAALLGVEGAIHAKYDAID
						KDTPIPTDRT/QLKWDIPRCHSV
						R*TVIHAPEGHAKFRRVLK\AW
			}			VVSHPDLVYWQGLDSLCAPFL
						YLNFNNEALVYACMSAFIPKYL
						YNFFLKDNSHVIQEYLTVFSQM
6628	26006	A	6670	1	500	IAFHDPELSNHLNEIGFIPD
6629	36996 36997	A	6678 6679	1	508 888	
6630	36998	A	6680	2	584	
6631	36998	A	6681	2	152	QVAKGMDALLQHLEDCGYR/S
0031	30999		0081	2	132	SKKKAQICRQQVTKWGDGSAL MGTSTTASL
6632	37000	Α	6682	733	894	
6633	37001	В	6683	1	1626	
6634	37002	Α	6684	1	1971	MAQVWANDNPPGLAVNQAPV
						LIDVKPGAQPIRQKQYPVPREA
	İ					LKGIQPGTKDYQPVQDLRLVN
						QATVTLHPTVPNPYTLLGLLLA
					EDSWFTCLDLKDAFFSIRLAPES	
			İ			QKLFAFQWEDPQSDLGCILLLY
						VDDLLLGHSTAVECAKGMDVL
						LQHLEDCRYKMSKKKAHICRQ
						QVRYLGFTIRKGERS\WEREKM
	İ					AVGVLTQTVGPWPRPVAYLSK
						QLD/EVSKGWPLCLRTLAATAL
						LAQEADKLTLGQNLNIKAPHA
						VVTLMTTEGHHWLTNARLTKY
						QSLPCENPHITIEVCNTLNPTTL
						LPVSESPGEHNCVEVLDSVYSS
		1				RPDLRDQPWASSVDWELYMD
		1				GSSFINSQGERCAGYAVVTLDA
						VIKAKLWLQGTSAQKAELIALT RAVELSEGQESLEELLGRYFYV
						SHLPAFAKAVAQLCITCRQHNA
						RQSPTVSPHIQAYGAAPFEDLQ
						VDFTEMPKCGGNKYLLVLTCT
						YSGWVEAYPTRTEKPYKKGKN
						DPSCTKGQCNPLELVITNPLNP
						HWKKGERVTLGIDRARLDPRV
			!			NILVRGGEVYERSPEPVFQTFY
			}			DELNVPVPEIPGKTRNLFLQLA
						ECVAQPLNVTSCYVCGGTVTG
						YQWPWKARELVPVDPVPDEFL
						AQKNYPDNFWVLKPSITGQYCI
6635	37003	A	6685	1	1461	
6636	37004	A	6686	1	1005	

SEQ ID NO:	SEQ ID NO: of peptide	Met hod	SEQ ID NO:	Nucleotide location of first	Nucleotide location of last codon for last amino acid	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
6637	37005	A	6687	1	174	
6638	37006	Α	6688	519	588	
6639 37007	A	6689	2	4527	GRAGLGWSSGGRGSRACSEGP AEARGREGDKGSAALSPPLPLT LGEEMAAERGARRLLSTPSFWL YCLLLLGRRAPGAAAARSGSAP	
						QSPGASIRTFTPFYFLVEPVDTL SVRGSSVILNCSAYSEPSPKIEW KKDGTFLNLVSDDRRQLLPDGS LFISNVVHSKHNKPDEGYYQCV ATVESLGTIISRTAKLIVAGLPR FTSQPEPSSVYAGNNAILNCEV NADLVPFVRWEQNRQPLLLDD RVIKLPSGMLVI
6640	37008	A	6690	1	879	
6641	37009	Ā	6691	407	594	
6642	37010	A	6692	1	3621	
6643	37011	В	6693	1	609	
6644	37012	В	6694	1335	1599	
6645	37013	A	6695	1	354	
6646	37014	A	6696	181	683	SRYFILLRKPTFPAMALLPVLFL VTVLLPSLPAEGKDPAFTALLT TQLQVQREIVNKHNELRKAVSP PASNMLKMEWSREVTTNAQR WANK/CTLQHSDPEDRKTSTRC GENLYMSSDPTSWSSAIQSWY DEILDFVYGVGPKS/PQYCPAG NNMNRKNTPYQQGTPCPG
6647	37015	Α	6697	199	969	
6648	37016	Α	6698	1	564	
6649	37017	Α	6699	153	427	
6650	37018	Α	6700	1	978	
6651	37019	Α	6701	3	419	
6652	37020	A	6702	9	255	VRAPAQGPDLAGGRRRCGSGA SCTPSPRGPASWSRSAAQVPRS SRWRAGSASS*N/GRQPAPPTSQ PPRAQPFAQPPGPWPLS
6653	37021	В	6703	96	283	
6654	37022	A	6704	172	365	PTYHLYPCHFWYQHRTRRIPRK SERLFPTPPPQLSWKGTWDY*I CLLCCNDSEGRDFLVLRVLK
6655	37023	Α	6705	3	1171	
6656	37024	Α	6706	2	740	
6657	37025	Α	6707	256	461	
6658	37026	Α	6708	1	414	
6659	37027	A	6709	1330	1386	YIDSILPK*NHDTLQSYCRYM\H SRQRAAS*PLSTGKLQQCLGLD LVLC

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
6660	37028	A	6710	207	1173	GGDRDRMTANHESYLLMASTO
0000	37028	^	0710	207	1173	NDMEDWVKSIRRVI\GDLSGGG
ŀ						IFGQKLEDTVRYEKRYGNRLAP
						MLVEQCVDFIRQRGLKEEGLFR
						LPGQANLVK\ELQDAFDCGEKP
						SFDSNTDVHTVASLLKLYLREL
				-		PEPVIPYAKYEDFLSCAKLLSKE
						EEAGVKELAKQVKSLPVVNYN
						LLKYICRFLDEVQSYSGVNKMS VQNLATVFGPNILRPKVEDPLTI
						MEGTVVVQQLMSVMISKHDCL
						FPKDAELQSKPQDGVSNNNEIQ
	1					KKATMGQLQNKENNNTKDSPS
						RQCSWDKSESP/TEKQHEQWIP
(((1	27020	 	(711	172	001	HSSIRQQNQQPKEQWSQAI
6661	37029	A	6711 6712	1605	1844	
6662 6663	37030 37031	A B	6713	123	699	
6664	37031	A	6714	123	462	
6665	37032	A	6715	1	2607	
6666	37034	A	6716	1	918	
6667	37035	A	6717	1	549	
6668	37036	A	6718	1	687	
6669	37037	A	6719	1	258	
6670	37038	A	6720	90	1621	HGLDLRTMNRSRQVTCVAWV
						RCGVAKETPDKVELSKGGVKR
						LIAEAEEKLQEEGGGSDEEKTG
						SPLEEG\MQSARTQARP\REPLE\
						DGDPED\DRTLDDDELAEYDLD
						KYDEEGDPDAETLG\ESLLGLT
						VYGSNDQDPYVTLKDTEQYER
						EDFLIKPSDNLIVCGRAEQDQC
						NLEVHVYNQEEDSFYVHHDILL
					}	SAYPLSVEWLNFDPSPDDSTGN
]						YIAVGNMTPVIEVWDLDIVDSL
						EPVFTLGSKLSKKKKKKKGKKSS
					j	SAEGHTDAVLDLSWNKLIRNV
		l				LASASADNTVILWDMSLGKPA
						ASLAVHTDKVQTLQFHPFEAQT
						LISGSYDKSVALYDCRSPDESH
				1		RMWRFSGQIERVTWNHFSPCH
						FLASTDDGFVYNLDARSDKPIF
		1			1	TLNAHNDEISGLDLSSQIKGCL
						VTASADKYVKIWDILGDRPSLV
]		HSRDMKMGVLFCSSCCPDLPFI
]		YAFGGQKEGLRVWDISTVSSV
						NEAFGRRERLVLGSARNSSISGP
						FGSRSSDTPMES
6671	37039	A	6721	1	549	1 0010001111100
00/1	13/039	ΙΔ.	10/21	<u> </u>	1379	l

SEQ ID	SEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first		*=Stop codon, /=possible nucleotide
ŀ	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
6672	37040	Α	6722	1	1318	AKQQLNLRTHMADENKNEYA
<u> </u>		ŀ				AQLQNFNGEQHKHFYVVIPQIY
		ŀ				KQLQEMDERRTIKLSECYRGFA
						DSERKVIPIISKCLEGMILAAKS
		ŀ				VDERRDSQMVVDSFKSGFEPPG
						DFPFEDYSQHIYRTISDGTISAS
•		ŀ				KQESGKMDAKTTVGKAKGKL
						WLFGKKPKPQSPPLTPTSLFTSS
ŀ						TPNGSQFLTFSIEPVHYCMNEIK
•		1				TGKPRIPSFRSLKRGWSVKMG/
		1				AALEDFSHLPPEQRRKKLQQRI
					DELNRELQKESDQKDALNKMK	
}		İ				DVYEKNPQMGDPGSLQPKLAE
ŀ		ļ				TMNNIDRLRMEIHKNEAWLSE
		İ				VEGKTGGRGDRRHSSDINHLVT
ĺ						QGRESPEGSYTDDANQEVRGPP
	1					QQHGHHNEFDDEFEDDDPLPAI
						GHCKAIYPFDGHNEGTLAMKE
		1				GEVLYIIEEDKGDGWTRARRQ
						NGEEGYVPTSYIDVTLEKNSKG
6673	37041	Α	6723	1	741	
6674	37042	Α	6724	1	1063	MPFYISDLSICGDRILRALCPQD
						LPTYSLHSRGKMRASCSRKFLD
						NNSSRLVSCNMGALISIWGTTT
	1					PPLHATILDSQPTVHPPLAKDCL
					•	PCGLQASASDLRARALQRLCQ
						QLPWVGSQPHTRSPSPQRGGKT
						GLFAGLASSVSMRPASPPSPAA
						DSCSACRFFARRPPLRVTWVKP
						SSALALCVSISDSIPGNLKALPA
						ETRAQLHHAEASLSQPPLQLRP
						FPKTSQAGDLQDLGPYVCVRK
						AVGKGDKQIRAVVKEHSVRSQ
]			1			ERIWHYPGITTANMPGHLGQNT
						ESGRDKLPMFGVWFPGCRFWG
1						LWVWRLP*LKLAAPCRPSRSLR
		<u> </u>				SSPISRRASTTRCLTVSGCPAAP
6675	37043	Α	6725	25	573	
6676	37044	Α	6726	1	483	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	•	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
6677	37045	ĪA	6727	1	996	MVGKAKWKPIELPLLRKIVNO
0077	3,043	1	0,2,			KQYHIPGGVAKISATIKDLKDV
						GVVIPTTSPFNSPIWPVQKTDGS
						WRKW\LS*QGQQFT/FTVLPHG
						YINSLALCHNLIRRDLDHFSLPQ
						DITLVHYIDDIVLTGSSEQEVAN
						TLDLLTSLDWALVETECLTMG
						HQVTMQPELTIMNWALSDPSN
						YKVGCAQQHSIIKWKWYIRDQ
						ARAGTTCKWTAAALQPLSRTS
						LKDSGERKSSQWAELRAVHLI
						MHFAWKEKWPDVQLYTDSWA
			i			VANGSAGWSGTWKNHDWKIA
						TTVIAQWAHEQSSHGSRNGGY
						TWAQQHGLPLTKADVATATDE
		1				CPICQQQRPTLSP
6678	37046	A	6728	2	1380	CLAGLFGFWSPCIPHLGVLLQSI
0078	37040	^	0720	-	1300	YLVTQKAA/SSEWGPEQEKALQ
				ł		EVQAAVQAALILEPYDPAGPVV
,						LEVSLADRDAVWSLWQAPIGE
						SQQRPLGFWSKALPSSA/DHKA
						CHAQQHSIIKWKWYIHDRARA
						GPEGTNSSARYAATM*KW\TAS
						ALQPLSRKSLKDSSEGKSSQWA
						ELRAVHLAVHVAWKEKWPDV
						RLDTDSWAVANGLARWSGTW
						KEHDRKIGDKEVWGRGTRIELS
						EWSKTVTIFVSH/VLLPRLPSIRG
						LTECLIHQHGIPHSIASDQGTHF
						TAKELQQWAHAHGIHWSYHVF
						HHPG\WGKVLQKAVYALNQRP
						IYEWKEESCLHTGVADALRGN
						WAEGHREHKALWLGLWSTWS
						QHPLRSLKTTRHHPGLGVLSED
						ICEAGGATEELSRASGFATGYG
						KRKEDTKKHKQHSVSDIIEQQH
						SLGLTEKTVKGTPTQGISMRPG
	İ					LYHKATEFQE
6679	37047	A	6729	1	2229	
6680	37048	A	6730	1	789	RDLQPFTSVTVHCRKGNDQTF
						GGPLDAGSELTLIPGDPKHHCG
						PPVKVGAYGGQVINGVLA\HPL
						IWLVQKTDGS/WRMTVDYCKL
						NOVVIPIAAAVSDVVSLLEQINT
						SPGTWYAAIDLANAFFSIPVHK
		I				AQQKQFAFSWQGQQYTFTVLP
						QWYINSPALCHNLIRRDLDCFS
						LPLDITLVHYIDDIMLIGSSEQE
			c .			VANTLDLFVRHLRARGWEINPT
						KIQGPSTSVKFLGFQWCGACQA
		1				IPSKMRDKLLHLVPPTTKKEAQ
	L	<u> </u>	L	L	L	II SKINIKUKELIEVEFT TKKEAQ

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
6681	37049	A	6731		553	MTVDYCKLNQVVIPIAAAVSD VVSLLEQINTSPGTWYAAIDLA NAFFSIPVHKAQQKQFAFSWQG QQYTFTVLPQWYINSPALCHNL IRRDLDCFSLPLDITLVHYIDDI MLIG\PRQLLACY/WALVETEHL TISHQVTMRPELPIMNWVLFDP SSHKVGCAQQHSIIKWKWYVH DWARAGPEGT
6682	37050	Α	6732	3	266	
6683	37051	Α	6733	3	582	
6684	37052	Α	6734	3	403	
6685	37053	Α	6735	1	2712	
6686	37054	A	6736	875	1506	LSIYLVTQKAA/SSEWGPEQEK ALQEVQAAVQAALILEPYDPA GPVVLEVSLADRDAVWSLWQ APIGESQQRPLGFWSKALPSSA/ DHKACHAQQHSIIKWKWYIHD RARAGPEGTNSSARYAATM*K W\TASALQPLSRKSLKDSSEGK SSQWAELRAVHLAVHVAWKE KWPDVRLDTDSWAV/ANGLAR WSGTWKEHDRKIGDKEVWGR GTRI
6687	37055	A	6737		633	MTVDYRKFNQVVTPMAA/AVP DAVSLLEQINTFPGTWYAAIDL ANAFFSIPVHEAHQKQFAF\LPQ GYINFPALCHNLIRRELDFFLLL QDITLVHYIDDILLIGSSEQEVV NTLDLLIHKRSKEAEHTAASRIR VSCLPEQKSHEQTLPWEQVPSS GDIKEYFPNAFVLLTTASLQGQ DNTSQLQLTWKAPEDIKMSKT DADADEEIEALRG
6688	37056	A	6738	2	1103	DLWPFTRVTLHRGKRNDQTFQ GLLDTGSELMLIPEDTKHHCGP PVKVEAYGGQVINGVLAQIQLT VGPVGSGTHPVVIYPVPECIIGI GILSSWQNPHIGSLTSRKTDGS WRMTVHYHKLNQMVTPIAAAI PDVVSLLEQVNTSPGSWYAAID LANAFFFIPVHKAHQKQFAFSW QGQQYTFTVLPQWYINSPALCH NLIRRDLDCFSLPLDITLVHYID DIMLIGSSEQEVANTLDLFVRH LRARGWEINPTKIQGPSTSVKFL GFQWCGACQAIPSKMRDKLLH LVPPTTKKEAQCL\QLLACY/W ALVETEHLTISHQVTMRPELPI MNWVLFDPSSHKVGCAQQHSII KWKWYVHDWARAGPEGT

of peptide sequence	hod	in USSN 09/540,217	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
		09/540,217		i	
37057		i	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
37057	•		sequence		
	Α	6739	11	2058	
37058	Α	6740	2	1173	WEVRQYTFTVVPQGHINSLVLC
					HDVVRRDLGHFFLPQDITLVHY
					IDDIMLIGSTPPITKKKAQHLVG
					LFGFWRQLVPHLGVLLWPIYQ
					VTQQAASFEWGPEKEKALQQV
					QAAVQAALPLGPYDPADPMVL
					EVSVADRDAVWSLWQDPIGES
		ļ			QWRSLGFWSKALPSSADNYSPF
					ERQLLACYWVLVETERL\QWV
					TPAEEDFNNQVDRMTHSVDTT
					QPLSPATHVITQWAHEQSGHG
					GRDGGYTRVQQHGLPLTKADL
					ATAGKVFQKAV*ALN*HSIYGT
					LSLIARIHRSRNQGVEVEVAPLT
					ISPSDPPAKFLLPVPPTLRSAGLE
					FLVPEEGMLPSGDTTVPLNWKL
					RLPPGHFGLLFPLHQQANKGVT
					VLVGVTDLDYQDEISLLPHNG
37059	Α	6741	823	2487	KELKLWKNRHKLLSYPTVGAA
					VTQLQNLTAMGVIGSHGARGQ
					VVALNRQRQGDLQPFTRVTVH
					WGKG/NMQIFGGLLDTGSELTL
		:			IPGDPKHHCGPPVKVGAFGG\Q
					VINGALARVQLIVGPVGPWAHP
					VVISPVPECIIDIDILSSWQNPHIS
					SLTGRVRAIMVGKAKWKSLEL
					PLPRKIVNQKQYHIPGGIVEISA
					TIKDLKDAGEVIPIPTTFPLNSPI
		1			WPVRKTDGCWRMTVDYCELN
		i e			QVVTPTAATVPDVVSLLEQINT
					SPGTWYADIDLANAVFSIPVHK
					AHQKQFAFSWQGQQYTFTVLP
					QEYINSLGLCHNLIQRDLDHFSL
					LQDITLVHYIEDIMLIGSTEQEIA
					NTLDLLFMAKEVWQWAHAHG
					IHWSYHVSHHPEAAGLIEWWK
					GLLKLQLQCQLGDNTLEGWGK
					VLQKTVYALNQHPIYGTVSPIA
					RIHRSRNKGVEVAPFTITPSDPL
]	PQFLLPVPATLCSAGLEVLAPE
					GGTLPPGDRTTIPLNWKLRLPP
					EDFGLLPSLSQQAKKGVTVLAG
					VIHPDYQDEISLLLHNGCKEDL
37060	A	6742	3	350	
			† <u></u>		
3	37059 37060 37061 37062	37060 A 37061 A	37060 A 6742 37061 A 6743	37060 A 6742 3 37061 A 6743 1	37060 A 6742 3 350 37061 A 6743 1 3339

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
6695	37063	A	6745	3	1377	AHLAADDFCTK*GWVSG\LPQG LDKLTLPGANLEMQPENLKED LVYLKKNHEELSMAIKRNESM AKGLQRALLQQQPEDDSKCFP RPQDLIRLYDIILQLTLEDREGK GLSFSFACVDKDVSVKMDTVP GVNLSCILNEMRDQDKKLVEK SCKDAQGWFFSLKALLEGSLVE TEVCYRTQLAQLQGLIRSMEQQ LCELCCDAEHQDHEHQVLLDV KTQLEQEIATYSRLLEVEDAQL ATQYSLSLASQPTREDLEKAIL KFIWNQNRAQIAKTILSKKSKA RGIMLPSFKIYYKTTVTKTAWC WYKNRHIDQWNRIENPEIRPHT YNHVIFDKPDKRKQWGKDSLF NKWCWENWLAICSKLTLDPFL NPYTKVNSRWIKDLNLQPKTIK ILKENLGNAIQDTGKGKDFMTK MPKAIATKAKIDKWDLIKLKSF
						CTALRNYQQSKPTTYRMGENF
6696	37064	A	6746	1	882	
6697	37065	A	6747	13	1402	STGSTHACDLLFSPSPVCLPPAA ATMTTSIR\QFTSS\SIKGSSGL GGG\SSRTSCRLSGGLG\AGSCR LGSAGGLGSTLGG\SSYSSCYSF GSGGGYGSSFGGVDGLLA\GGE KA\TMQNL\ND\RLGSYLDKVPC PWKEANTELEVKI\RDLVTRGR APGPARDYSQYYRTIEELQNKI LTATVDNANILLQIDNARLAAD DFRTKFETEQALRLSVEADING LRRVLDELTLARADLEMQIENL KEELAYLKKNHEEEMNALRGQ VGGEINVEMDAAPGVDLSRILN EMRDQYEKMAEKNRKDAEDW FFSKTEELNREVATNSELVQSG KSEISELRRTMQALEIELQSQLS MKASLEGNLAETENRYCVQLS QIQGLIGSVEEQLAQLRCEMEQ QNQEYKILLDVKTRLEQEIATY RRLLEGEDAHLTQYKKEPVTTR QVRTIVEEVQDGKVISSREQVH
6698	37066	A	6748	2	452	FLLPSSFSCTHSLLSPTITIVPGP DFNLASHIIPDTTPDPHDCISMIH LTFTPFPHISFFPVPHPDHA*FID GSSTRPNRHTPAKAGYAIAQAT IPPLRTSHFLSIVEIYPQGNNFSV FHVLFYYSSGIIQAPSLPYTLRR DRRPGTSRA
6699	37067	Α	6749		1791	DIGG OTOKA

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
6700	37068	A	6750	1	219	
6701	37069	A	6751	2	528	TLTVESKSIKAYKLSLQFPHFTC PKTGHALQVSSGFVESPTVTIVP GLDINLASHTIPDNTPEPHDCIS LIHLTFTPFPHVSFFPVPHPDHT WFIDGSSTRPNCHSPAKTGYAI VSSTLIIDATALPPSTTSQQAELI ALTRALTLAKGLPINIHTDSKY AFHILHHHAVI*AER
6702	37070	В	6752	1	1744	
6703	37071	A	6753		1376	MCKRMPGLQASQTVCPFYDKN YLDLKRKGTEIKRRERLKCGTK IERRKRLRDSERLEKRIKRGCLP DLKLSTEIQINRLRGLYTLLRKE RRFGLQETINYAEVTGIYVKELI EDREHLNSGKYIFNIFDIKVPRV VFGIIDPYMLWEKKEGTKEGIK EGMKRESQKRIKWERSFITFIAL NTADRTLMSSAGCYRKQVKIHI DALPPPNSQTLKKRGGASLIISF RRQSGPSAAGLWEFARGPFQTL FAWLSPAEAAEQQRLLPASSAE SFIAEGDPPDAGQSSPPSSLHAA AATALMLLEALKITNYAQLTLY SSPNFQNLFSSSHLTHILSADWL LQLYSLFVESPTITIVPGPDFNPA FHFIPDTTPDPHDCISLIHLTFTP FPHISFFPVPHPDHTWFIDGSST RPNRHTPAKAGYA/DGRNTSYC VLSPVGNSPLVPNPGSARCTEG
6704	37072	В	6754	400	820	GLLYARA
6705	37072	В	6755	1	1190	
6706	37074	A	6756	1	1002	
6707	37075	В	6757	1	3216	
6708	37076	A	6758	1	4020	
6709	37077	A	6759	1	1155	
6710	37078	A	6760	2	346	SSYLMHILSAPNLLQLYSLFVES PTITIVPGPDFNLASHIILDTTPD PHDYLSDPPDIHS/TFP/DISFFPV PHPGHTWFNDGSSTRPNRHSPA KAGHAIVFSTSIIEATALPPSTT
6711	37079	Α	6761	189	1201	
6712	37080	Α	6762	1	231	
6713	37081	Α	6763	1	834	
6714	37082	A	6764	470	479	PRNPSSFLQVQVQHRFLQLFVP FHFVFFLASLYVMVTLTTWFR* DPCVSGMALNCSFLKIASSSGE YLRPTPSYLNQVVRVTMT
6715	37083	Α	6765	21	706	
6716	37084	Α	6766	1	158	

SEQ ID	ISEO ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence		in USSN 09/540,217	location of first codon for peptide sequence		*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
6717	37085	С	6767	78	401	
6718	37086	Α	6768	536	694	
6719	37087	C	6769	1	1410	
6720	37088	В	6770	1	1836	
6721 3708	37089	Α	6771	306	487	VGFLGILF/CFEAIVNVSSLMIW
		1				LSVCLLSVYKNACDFCTLILYP
						ETLLKLLISSSSLYLW _
6722	37090	Α	6772	714	1096	FFLPMSMECSSICLYPL/CISLRS
						GL*FSLKRSFTSLVSWIPRYFILC
						KSQGMRWLGLGSEA*YSCLLIL
						IRKITQNSVEVLGRRKFLGGGM
:						EREWVMFLRAASSGIRGSVGT
			(===		2506	NFKSESRHQISCASV
6723	37091	C	6773	1	2796	
6724	37092	A	6774	126	490	FFLPMSMECSSICLYPLLFPGAV
				Ε		VCSSP*RGPSHPL*VGFLGILF/C
					}	FEAIGNGSSLMIWLSVCLLLVY
						KNACDFCTLILYPETLLKLLISL
						RRFWAETMGFSRYIIMSSANRD
6725	127002	 	(775	t	550	NLTSSFPN
6726	37093 37094	A	6775 6776	578	865	MCPRDYGMLCLCSH\CFKEHLY
0/20	37094	A	0770	378	803	FCLHFVMYPVVIQEQVVQFPCS
						*AVLSEFLNPEF*FDCTVV*ETV
						CYNF*SFTFAEKSFTSNSVVNFG
						IGVVWC
6727	37095	A	6777	1	2091	
6728	37096	С	6778	1	1458	
6729	37097	В	6779	1	504	
6730	37098	Α	6780	70	511	NHASPGIRNLFLHPRGLRAITIA
						VFCKQNTYIRLEPFKINVLEQIT
						KHIEKLQCGGVVKQLSRRGNN
						QHISSTYDINRADTQVRRAVNN
						YDII\A*ATVSMASRSIRCGSVGS
ŀ						LPSSNLPINSVSIRCRARQEGIM
6721	27000	ļ	£701	2	306	SSPGQQVGFIA
6731 6732	37099 37100	A A	6781 6782	3	3567	MHIVVETALSASWQNKAKPPA
0732	13/100	^	0782		3307	RVLLQVVPNVWFLVAVVWEL
						YPSLDLMDRSIECSSSPATEQS
		i				WTENDYDKLREEGFRRSNYSE
		ĺ				LQEDIQTKGKDVENFEKNLEEC
			ļ			ITRITNTQKCLKELMELKTKAR
						ELREECRSLRSRCDQLEERVSV
						MEDEMNEMKQEGKFREKRIKR
						NEQSLQEIWDYVKRPNLRLIGV
						PESDGENGTKLKNTLQDIIQENF
		}				PNLARQAKVQIQEIQRMPQRYS
						LTRATPRHIIVRFTKVE
6733	37101	A	6783	386	550	
6734	37102	A	6784	1	594	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
6735	37103	l _A	6785	<u> </u>	11324	MGKKQSRKTGNSKRQSASPPP
0,55						KERSSSPAMEQSWMENDFDEL
						REEGFRRTNYSELQEGIQTKGK
						EVKNFEKNLDECITRITNTEKCL
						KELMELKAKARELREECRSLRS
					:	RCDQLEERHHTTPIPKLTTYLE
						VKLSPANVKEQVITNCLSDHSA
						IKLELRIKKPTQNPSTTWKLNN
						LLLNDYWVNKEMKSEIKMFFE
						TNENKNTTYQNLWDAFKAVCR
						GKFIALNAHKGKQERSKIDTLT
						SQLRELEKQEQTHSKVSRRQEI
						TKIRAEPKEIETQKTLQKINESR
						SWFFERINKIDRRLARLIKKKRE
						KNQRDAIKNDEGDITTDPTEIQT
					:	TIREYYKHLYANKLENLKEMD
			ŀ			KFLNTYTLPRLNQEEAESLNRPI
						AGSEIVAIINSLPTKKSPGPDGFT
						AEFYQRYKEKLVFGAGYFGM
						WALAALPSNLLKLSQLCQEAA
						EVNVLVQFVCIC*SCLNSVRRQ
6736	37104	C	6786	1	1719	
6737	37105	В	6787	1	1098	
6738	37106	C	6788	1	1782	
6739	37107	A	6789	3095	3217	CRCLQMAM*WS*AIALSMKIS
(7.40	27100	ļ. —	6700	,	5264	MSPRKPEV*IKVLAGLCCL
6740 6741	37108	A	6790	1	5364 2505	
6742	37109 37110	C	6791 6792	37	1214	KLLQDLPFSINSSAVEKISMSTT
0742	3/110	^	0792	37	1214	GQVIRCKAAILWKPGAPFSIEEV
						EVAPPKAKEVRIKVVATGLCGT
						EMKVLGSKHLDLLYPTILGHEG
						AGIVESIGEGVSTVKPGDKVITL
						FLPOCGECTSCLNSEGNFCIOFK
						OSKTOLMSDGTSRFTCKGKSIY
						HFGNTSTFCEYTVIKEISVAKID
	,					AVAPLEKVCLISCGFSTGFGAAI
						NTAKVTPGSTCAVFGLG\GVGL
						SVVMGCKAAGAARD/IIGVDVQ
						QGRKF*GRAQELGATEC\LNP\Q
						ALKKPHFKEVLFDMTDAGIDFC
						FEAIGNLDVLAAALASCNESYG
						VCVVVGVLPASVQLKISGQLFF
						SGRSLKGSVFGGWKSRQHIPKL
						VADYMAEKLNLDPLITHTLNL
[-	DKINEAVELMKTGKW
6743	37111	В	6793	1	2280_	
6744	37112	Α	6794	1498	1764	LVYLKVMGRMEPSWKTLCRIL
	}					SRRTSANLSKAGQCSDSGNTEN
						TTKILLKKSNSKTHNCQIHQS*N
1						EGKNVKGSQRERSGYPQRGRPS

	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
6745	37113	A	6795	3	640	
6746	37114	Α	6796	1	996	
6747	37115	Α	6797	1	933	
6748	37116	Α	6798	74	1128	LLLDL*VD*SPSRPLEMRFRV*P
						MKRCTTLKKNYLSFLVYISRNL
						*NRVKAIM/VGKAEWKTIELPL
						PRKTVNQKQYRIPGGIAEISTTI
						KELNNAGMVIPTTSPFNSPIWP
					•	VQKIDGSWRMTVDYCKLNQV
						VTPIAATVPYVVSSLEQIN/TSPG
					•	T/WNRKRFCNRSRLLHKLLCHL
						GHMTQQI*CFLRCQWHIGMLF
		1				GPEICGCTSLSGQKL/WKIFVSH
						AASFEWSPEQEKALQQVQAVV
					QAALPLGPYDPADPVLLEMSV	
						ADRDAVWILWQAPISESQWRPP
	}				GFWSKALPSSADNYSPFERQLL	
						ACYWTLVETERLTMGHQVTIPP
						ELPIMNWVFSDLSSPKMCHPQQ
6749	37117	Α	6799	1	229	
6750	37118	Α	6800	1	1011	MVSTPATLPSLPKPALMASWG
						VPYDQLTKEEKTRVWFTDGSA
					RYAGTTQKWTAVALQPLSRTS	
						LKDSSEGKSSQWAELQAVYLV
						VHFAWKEKWPDVGLYTDSWA
						VANGLAGWSETWEKQDWKIG
						DKEIWGRGMWMDLSEWSKAV
					:	KIFVSHVSÅHQRVTSAEEEFNN
						QVDRPL/PVFTQWAHEQSGHSG
						RDGGYSWAQQT\GLPFTKADL
						AMATAECPICQQQRPTLSPLYS
			,			TIPQGDQPATWWQIDYIGPLPS
						WKGQKFVLTVIDTYSRYRFAY
						PAHNASAKTTIHGLIECLIHCYG IPHSIASDOSIHF/TTKEVQ*WAH
		1				AHGIHWSYHVSHHPEAAGL
6751	37119	A	6801	2	739	HKMGHAQQHSIIKWK*YICDW
0/31	3/119	^	0001	2	139	ARAGPKGTT\APMASWGVLYD
						QLTEEEKTRAWFTDG\$ARYAG
						TTQKWTAAALQPLSRTSLKGS
	1					GEGKSSQWAELQAVHLVVHFS
				1		WKDKWPDVRLYIDSWAVANG
						LAGWSGTWKKHDWKIGDKEI
				1		WGRGMWMDLSEWPKPVKIFG
				1		SHVSAHQWVISAEEDFNNQVD
				1		KMTCSVDITQPLSPATPVITQW
						AHKQSGHGGRDGGYTWAQQH
						GLPLTKTGLAMATAECPI
6752	37120	A	6802	1	354	O. D. H. I O. M. H. I I I I I I I I I I I I I I I I I I
6753	37121	A	6803	3	592	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	1	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
6754	37122	IA	6804	126	365	QLAEPPHWLPD*KGAGFEWGP
					<u> </u>	EQKKALQ*VQAAVQAALPLGP
	ļ		}			YDPADPMVLVVSVADKDAVWI
						FHLSGSDRWRRTYRCL
6755	37123	A	6805	1	636	MEFNDSMHNTFDHIWRTKEYD
0733	37123		10003	i	050	EAGWLLLSSLDKAMKENDELR
						DSNSQIQKQILSLQSSKIILNLTE
						GTVVTQLQNLNSVGIIGSQGGR
			1			GQVAALSLQRQGASFEWSPEQ
		İ				
						EKALQQVQTAVQAALPLGPYD
		1				PSDPMVFEVSVADGNAVWSLW
		1				QAPIGELQWRTLEFWSKALPSS
		1				ADNYSPFGRQLLACYWALVET
		ļ				EHLTVDHQVT/M*PELPIM
6756	37124	Α	6806	1	672	
6757	37125	A	6807	308	470	
6758	37126	Α	6808	1	1908	
6759	37127	A	6809	1	1218	MGIIGSRGGRGQVAALKRQRQ
		1				VTKKAASFEWGPEEEKALQQV
						QAAVQAALPLGPYDPADPIVLE
						VSVADRGAVWSLWQVPIGESQ
						QRPLGFWSKALPSSADNDSPFE
						RQLLACYCALVETERLDMGHQ
						VTMRPEVLIINCVLSDPSSH/Q\G
						WRLCMDSATWTSTYQ/A*PGY
						GHC*VPNLPAAETNTDPSIWHH
						S/YG*LASYLVIG*LHWTS\$IME
						GAEVCPHWNRHLLRI*VCLSCT
						QCFCED/SHPWIHE/ASYLPSWH
						ST*HCL*SRHSPYS*RSVAVGSC
	ļ.					SWNSLVLPCSPSS*SSWINRMV
	:					EWPFEVTITMPTR*QYSAGLGK
	:					SSSEGCVCSESASNIWYCFSRSQ
						DSWVQESRGGSGSGTIHHHF**
						PTSKIFASCFLDIMFCWT*GCS/S
	ľ					*WRNAATRRHNNNFI
6760	37128	Α	6810	901	1142	GPEDLQGLFILSFQDPLL/SHPV
0,00	37120		00.0			CLEALAPFPSPPVVPISSFPPHVA
						GEFWLYPSGAA/RASSDAQSWT
						VIAVSPLRGLNSWG
6761	37129	В	6811	1	1452	VIAVSI ERGENSWG
6762	37130	A	6812	122	1838	
6763	37131	A	6813	101	249	MTHVTALRTSSNLMRN*RKVG
"	31131	``	10015			EAKSQQEH*S*QACHLKSRMSS
						SRLWIN
6764	37132	A	6814	1	1074	JIC WIIV
6765	37132	A	6815	310	397	LTIWQDSNPGSCL*GCSVCAHA
] , 03	15,155	1.,				/CVGGQHMVCF*PSYSLYLKSE
		1				PKEGTGRNKDPDLNSQNRKGF
						LEQIHFQ
6766	37134	A	6816	1	583	DEQUIT Q
0700	13/134	ΙΛ.	10010	Ι'	505	

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
6767	37135	Α	6817	1	486	
6768	37136	Α	6818	2	1001	
6769	37137	Α	6819	1	2307	
6770	37138	Α	6820	3	393	
6771	37139	Α	6821	1	1776	
6772	37140	Α	6822	1	1140	
6773	37141	Α	6823	1	558	
6775	37143	A	6824	3	206	PRVRIFFSITWYSLRVLTEISLW GGLILVVIITIQYDMTRTRNKYL HTNSLVALANMSAQFR\SLHQY AA\QRIIRLFSLLSKKHNKVLEQ ATQSLRGSLSSNDVPLPDYAQD LNVIEEVIRMML\EIINSCL\TNS LHHNPNLGITPWLLQ\RRSLWN NFR\THPSFQGINGKILIWVIF/LS /SRLQGLAGKLGAELSVERVLEI IKQGVR\SLPKDRLKKFPELKFK YVEEEQPEEFFIPYVWSLVYNF SSRPVLGIPQDIPAVSPWVSD PSASRPAWHAPALHPSPVPARG WQPPTGLHPLREQRGLGQHM* QSSGQQPQGGVGPGSHFPGAL WAQL
6776	37144	A	6826	200	601	EMGSCDQWQRPCVPWEAVVM PRDP/LKMALAVVWPKMKAPR VTKRTP*LTSTHQLAGLTSSISE SLA*MRHSPGKTKMKQAARPP MTEITLPMSGMKS/VQQQGEQE PDQRLQHSPPPLPPHVLLHWHP LVAQPQA
6777	37145	В	6827	25	1407	
6778	37146	A	6828	1786	1795	HC*LNALPSGETWGQKRGLPG TQLPLPL/VPLRAPKGRSPKGSK GPTHHTWALSSFSLTPSQVTVG TEDQATSASGRPWLRIGTGPGG GRVWAGRDS\RGVPALTVLHV LSSLGRETP*RPCLRGPTAPAGP KRAPGKWLPWSQAPPCGCCPR *LSHVL*DRNRCAW*KGGGRG L/LRAPTPVPELAPHPCGPAPTL SKPLSSPNMPAQGLTQE*PGSV ANLPPYGPALLCLGRREWDER GQVG*VPTESEME*RRTEPLKT PGDCSAAASKIPSFFSNRTPFPPL CPHKNKLKGT
6779	37147	Α	6829	1	311	· · · · · · · · · · · · · · · · · · ·
6780	37148	Α	6830	1	402	
6781	37149	Α	6831	1	750	
6782	37150	С	6832	130	450	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
	ļ			sequence		
6783	37151	Α	6833	1	875	MRLLRLKLKVEQHVELYQKYS
]			NNSWRYLSNRLLAPSDSPEWLS
						FDVTGVVRQW/VWSRGGEIEGF
						RLSA\HCSCDSRDNTLQVDINGF
		İ		,		TTGRRGDLATIHGMNRPFLLFM
						ATPLERAQHLQSSRHRRALDTK
						YCFS\STEKNCCVRQLYIDFRKD
						LGWK\WIHE\PKGYHANFCLGP
		į				CPYIWSLDTQYSKVLALYNQH
						NPGASAAPCCVPQALEPLPIVY
						YVGRKPKVEQLSNMIVRSCKC
						NLGPAPPRPAPAGPAPPRPAPV
		İ				ALPMGAVFKDTRAPSPPGAPLK
		<u>L</u> .				MERGKKKKKK
6784	37152	В	6834	162	635	
6785	37153	Α	6835	2	1846	
6786	37154	Α	6836	322	2104	
6787	37155	A	6837	1	352	
6788	37156	В	6838	92	1774	
6789	37157	A	6839	260	1016	DALLIN CAN ADDRIVE DELLA MANAGEMENTA
6790	37158	Α	6840	479	1937	DVEHVGKMRRRWRRRVVVAD
1						GNVVKPMSCAGDLQ/PFTRVTV
						HWGKGNDQTFQDLLDTGSELT
					}	LIPGDPKRHCGPPVKIGAYGGQI
						INGVLAQVQLTVDAVGPWTHP
						VVIFPVPECIIGIDMLSSRQNPHT GSLTGRVWTIMVRKAKWKPLE
						1
						LPLPRKIVNQKQYHIPEGIVEIS ATIKDLKDAGVVIPTTSPFNSPI
						WPVQKTDGSWRMTVGYCKLN
						QVVTPIAAAVPDVVSLLEQINT PPGTWYAAIDLANDF
6791	37159	В	6841	20	26	PPGT W TAAIDLANDF
6792	37160	В	6842	2	207	
6793	37161	В	6843	187	405	
6794	37162	A	6844	451	770	LFLFLLSSHPRRSSASWYRHRH
07,54	37102		10044	1	1,10	PHHPSAARLPVADSASSSSSPSS
]	SSPSPSSSSSSSSSPPSSSSFVYPHC
					:	QPPAPHFPH*HSR*QGRLFHFLP
						LPQLPSSPLSPWW
6795	37163	A	6845	1	417	- C
6796	37164	A	6846	191	314	
6797	37165	A	6847	129	212	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
6798	37166	A	6848	1	192	CALNAGL/IGENEAKKRRSKVT
						QEADFPGSMDGASNFVRGDAI
						AGILIMVINIVGGLLVGVLQHG
		l				MSMGHAAESYALLTIGDGLVA
		l				QIPALVISTAAGVIVTRVSTDQD
	ŀ					VGEQMVNQLFSNPSVMLLSAA
		l			·	VLGLLGLVPGMPSLVFLLFTAG
		l				LLGLAWWIRGREQKAPAEPKP
						VKMAENNTVVEATWNDVQLE
						DSLGMATGLLT*PIFRAQWTGQ
		l				VTLFAAMPSPGSSSWSLTLSAG
						CWSACCNMA
6799	37167	В	6849	65	1129	
6800	37168	В	6850	46	1038	
6801	37169	Α	6851	2	413	PSFQGPVSLPSITVVSIDSQASKP
		l				LKTPQLWCQLRQYSFKHSFLV
		i				VPTCPVPLLG*DTLTKLSASLTI
		l				PGLQPHLIATHLNPQVEDTSIPS
		1				LATYHAPLTISLKPNHPYPSQC
						QYPIPQHALKGLKPVITHLLQH
6802	37170	С	6852	35	346	
6803	37171	С	6853	239	484	
6804	37172	Α	6854	1	1020	
6805	37173	Α	6855	3	131	
6806	37174	A	6856	1	956	MAGALPPASLPPCSLISDCCASN
						ERSSMGIGPSEPCAGYNLLVCR
		1			1	LTKKKRIRSTNKVWDYVKRPN
						LRIIGVPEEEENSKSCENILGEIIE
						ENFPSLATDLDIQIQEAQRTPQK
		l				FITERSLPRYIVIRLSKVKTKERI
		l				LRAVRQKHQIFLTQRLKTNTAQ
			ĺ			SPQKLPGPSQTLLVTLTREITIVP
		l				GLDFNPASHIIADTTPDPHDCIS
						LIHLTFIQFPHISFFPVPHPDHTW
						FINGSSTRPNRHSPAKAGYAIVS
		l				STSIIEATTLPSSTTSQQAEFIAL
		l				TQALILAKGLCINIYADSKY/AF
						HILHHHAVIWAERDTFTG
6807	37175	Α	6857	1	1269	
6808	37176	Α	6858	1	741	
6809	37177	В	6859	411	1257	

SEQ ID		Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide	hođ	in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
6810	37178	Ā	6860	438	3997	SPLPHSTPISHPAACFKKIKACY
						HLPATA WPFKAYKLSLQFPHFT
						CPKTRQGLQHNSHKNTRALPD
						YRLRLISQTPIPSTKQQLLSFLG
ł						MVSKVRILTQELGPRPIAFLSKQ
						LDLTVLTQPSCLHAAAAAALIL
	1					LKALKITKYAQLTLYSSHNFQN
						LFSSSYLMHILSAPWLLQLYSLF
						VESPTITIVPGTDFNPASHIILDT
			İ			TPDPHDCISLIHLTFTPFPRISFFP
		İ				VPHPNHIWFIDGSSTRPKCHSPA
			1			KAGY
6811	37179	Α	6861	245	505	
6812	37180	Α	6862	346	522	PAPEFWVHG*NVSPLSL*SKEQ
		1				EDRGFISQGRSPDLSHGTKFHV
						RPCEETTKQALCEQH
6813	37181	Α	6863	3	879	SGDLPWEINPPSSYTLLCEKDPP
						TTSGPQTNQPKKH\LTNFKSETK
						ETRFIRGPTTPAPVTDWEGSLPL
1						VFNHSRDTSLIIHPGFRGVRPRR
		1				DACLGPSPLAASPTFLGKGPAA
	ł					PRQTELGPNSSSASAPPPYNPFI
		1				ASPPHTWSGLQFPSMTSPPPPA
						QQFTLKKVAGAKGIVKDLINLT
						FKVYNNRKKLQFLASTVRQTP
						ATSPAHKNFQTPELQQPGVPPE
						PPPRGACYKFQKSGHRAKECLQ
						PRIPPKPHPICVGPHWKSDCPTH
		ŀ		i		LAATPRAPGTLAQGSLTPSQIFL
6814	37182	Α	6864	143	331	DTSTPSLVTDHASLTISLKPNHF
		ŀ				YPD*CQYPIPQHALKGLKPVITC
						LLQHGLLKPINSPYPRA
6815	37183	A	6865	1	783	
6816	37184	Α	6866	32	415	WRHALKGLKPVITRLLQHGLL
						RPINSPYNSPFLLVLEPDKPYRK
						RK*ISPPNRHILASVEVLKNQVN
						SHEFERSALIHTLTSGLAPCSLH
						KMCSKLPRSFLSVGSGTGKEAR
						SKILTFRTASRHLVL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
İ				sequence		
	<u> </u>	<u> </u>				
6817	37185	Α	6867	1	1374	MSELPFTIASKRIKYLGIQLTRD
				1		VKDLFKENYKPLLKEIKEDTNK
				1		WKNIPCSWVGRINIVKMAILPK
				1		VIYRFNAIPIKLPMTFFTELEKTT
ŀ]		LKFIWNQKRARIAKSILSQKNK
	1					AGGITLPDFKLYYKPTETKTAW
						YWYLNRDIDQWNRTEPSEITPH
	•					TYNYLIFDKPEKNKQWGKDSL
	ļ					FNKWCWENWLAICRKLKLDPF
			•			FTPYTKINSRWIKDLNVRPETIK
						TLEENLGITIQDIGVGKDFMSKT
		1				PKAMATKDKTGKWDLIKLKSF
						CTAKETTIRVNRQRVAALLNSA
					}	VFQNAVEVVVWPSSEVHVKFL
	1					FCKIISCAGQKQPIKFDGSSSLS
						GVATAMMSATIAHLKRIKACY
						RSPVTAWTFKAYKLSLQFPHFT
						CPKTGQALQ*HT*KGLKPVIAH
						LLQHGLLKPINSPYNSLILPVQK
						PDKPYRKLESFTSKAIKWHQIPS
						\PGKHLC**GS*RSPVTAWTFKA
		İ				YKLSLQFPHFTCPKTGQALQEA
						GVIHFKGHQMASDPITQGNTYA
						* · · · · · · · · · · · · · · · · · · ·
1			<u> </u>			NKVAKEAASVLTSVPHGQFFSF
		1	ĺ	1		SSVTPTYFPTEIDFTYMPQVRKL
		<u> </u>		100		KHLMV
6818	37186	A	6868	122	281	
6819	37187	Α	6869	2	2144	
6820	37188	Α	6870	224	418	LSQWRHALKGLKSVITRLL*HG
		1	ļ			LLKPINSPYNSPIVPVLKPDKTY
						RKVESFTAKTIKRCQIPLL
6821	37189	Α	6871	5	289	
6822	37190	Α	6872	1	2205	
6823	37191	C	6873	1	2082	
6824	37192	В	6874	1	1614	
6825	37193	Α	6875	1	2067	
6826	37194	A	6876	1	1152	
6827	37195	Α	6877	1	798	
6828	37196	С	6878	488	788	
6829	37197	Ā	6879	90	765	NTVFGLVVLPLELKLRIFRLLD
	1		1			VRSVLSLSAVCRDLFTASNDPL
		1	1			LWRFLYLRDFRGDFRNDIYSWT
		1	<u> </u>			TKDFEYYINLVDKTEARFEKVD
		1				1
		1	ł			SDFESFTVGKMPSNSIACTEKSF
			1			MKGRVSQCDKKPVITRLLQHG
						LLKPINSPYNSPILPVLKPDKPY
						KLVQDLHRINQIVLPIHPCGAQP
						HTLFCPQYLPPQLTIPVPDLKDA
1	I	1	1	1	1	FFTIPLQPSSQ\PLFAFT*TDPDT

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
6830	37198	A	6880	1824	2286	PRVLRPNSPRNISPISNRIFSD*R CPIASEASWTITDALG/RLLQWR HGLLNPINSPYNSPILPVQKPDK PYRLVQALLSLPQLPLFLAWTSI WPPTLFRITLLNPMTVSL*ST*H
6831	37199	A	6881	930	1289	SLHFPMFPSFLFLTLITLGLLMA VPLGLIATHQQRRAML LSQWTDGLLKPMNSPYNSPILP
0631	37177		0001	250		VLKPDKPYK*VQDLRLINQIVL PIHPVV\PNPYTLLSSIPPSTTHY SVLDLKLAFFTIPLHPSSQPFFAF TWTDPDTHQAQQIT*AVLPQSF SDSPHE
6832	37200	Α	6882	2	196	IHGCKFIQCKRILSNFTEVADHV HDQIRL*TGIRLLTQPQPRLSCR HQVAQHSVCRYLCQLPLL
6833	37201	A	6883	2	1441	LNQEEGESLNRPITGSEIEAIINS LPTKKSPGPDGITAEFYQRYKE ELVPFLVKLFQSTEKEGILPNSF YEASIILIPKPGRDTTKKENFRPI SLMNIDAKILNKILANRIQQHIK KLIHNGQHINRTNDKSHIIISIDA EKTFGKIQQPFMLKTLNKLGID GTYLKIIRAIYDKPTANIILTGQ KLEAFPMKTGTREGCPLSPLLF NIVLEVLARAIRQEKEIKGIQLG KEEVKLSLFADDMIVYLENPIV SAQNLLRLISTFSKVSGYKINVQ KSQAFLYTNNRQTESQILSELPF TIASKRIKYLGIQLTRDVKDLFK ENYKALLNEIKEDINKWKNIPC SWVGRINIVKMAILPKVIYRFN AIPIKLPMTFFTELEKTTLKFIW NQKSARIAKSILSQKNKAGGIM LPDFKLYYKATVTKTAWYWY QNRDIDQWNTTEPSEIMLHIYN Y/LIFNKPEKNKKWGKDSLFNK WCWENWL

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
				Sequence		
6834	37202	Α	6884	69	2415	DKTSRGTIRQQRLRFTNIRCSAA
		ŀ				TAADIQANRVWRGPPANSNRP
	1					AA\RVLTVRRKTNKQKGHPHQ
						KPICTSPSSNTKEIQTTIREYYKH
						LYANKLENAEERDKFLNIYTLP
		1				RLNQKEVESLNRPITGSEIEAIIK
			ŀ			SLPTKKSPGPDGFTAKFYQRYK
	1					EELVPFLLKLFQSIEKEGILPNSF
						YEASIILIPKPGRDTTKKENFRPI
}	1					SLMNIDARILNKILANLIQQHIK
}						KLIHHDQVGFIPGMQGWFNICK
; 						SINIIQHMNRTKDKNHMIISIDA
						EKPFDKIQQPFMLKTLNKLDID
		ŀ				GTYLKIVRAKYDKPTANIILNG
						QKLEAFPLKTGTRQGCPLSPLL
						FNIVLEVLAREIRQEKEIKGIQL
ļ	ŀ					GKEEVKLSLFVDDMIIYLENPIV
		ŀ				SAQNLLKLISNFSKVSGYKISVQ
		ľ				KSQAFLYTNNRQTESQTMSELP
						FTIASKRIKYLGIQLTRDVKDRF
		ŀ				KENYKSLLSEIKEDTNKWKNIP
		ŀ				CSWVGRINIVKMAILPKVIYRF
		ŀ				NAIPIKLPMTFFTELEKTTLKFIR
						NQKRARIAKSILSQKNKAGGIT
						LLDFKLYYKATVTKTAWYWY
						QNRHVDQWNRTEPSEIMLHIY
						NYLIFEKPDINKQWGKDSLFNK
		ŀ				WCWENWLAICRKLKLDPFLTS
		}				YTKINSRWIKDLNVRPKTIKTLE
						ENLGNTIQDIGMGKDFMSKTPK
						AMATKAKLDKWDLIKLKSFCT
		}				AKETTIRVNRQPTGWEKIFATY
6835	37203	A	6885	1	2358	
6836	37204	В	6886	1	1443	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
6027	127205	A	6887	453	2913	KMGLAREWILRFLVLILQEIRS
6837	37205	A	0887	452	2913	1
						MRQKENKDIQDLNSALHQADL
						IDIYRTLHPKSTEYTFFSAPHCI/
						RTYSKIDHTLGSKALLSKCKITE
ĺ						IITDSQTTVLSELRIKKLTQ\NRS
ĺ						ATWKLN/MLLNDYWVHNEMK
						AEIKMFFETNENKDTTYQNLW
ĺ						DTFKAVCRGKFIALNAHKRKQ
ĺ		ŀ				ERSKIDTLMSQLKELEKQEQTH
İ						SKVSRRPIPSSEIEAIINSLPTKKS
İ	Į					PGPDGFTAEFYQRYKEELVPFL
						LKLFQSIEKEGILPNSFYEASIILI
						PKPGRDTTEKENFRPISLMNIDA
						KILNKILANRIQQHIKKLLHHDQ
						VGFIPGMQGWFNICKSINVIQHI
ļ.						NRTSDKNHTIISIDAEKAFNKIQ
						QLFMLKTLNKLGTNGMYLKIV
						RAIYDKPTANIILNGQKLEAFPL
						KTGTRQGCPLSPLLFNIVLEVLA
		ļ				RAIRQEKEIQGIQLGKEEVKLSL
		İ	l			FADDMIVYLENPIVSAQNLLKLI
		1				SNFSKVSGYKINLQKSQAFLYT
						NNRQTESQIMSELPFTTASKRIK
						YLGIQLTRDVKDLFKENYKPLL
						NEIKEDTNKWRNIPCSWVGRIN
						IVKMAILPKVTYRFNAIPIKLPM
						TFFTELEKTILKFRWNQKRAHI
						AKTILSQKNKAGGIRLPDFKLF
1		{				YKATVTKTARYWYENRDIDQ
					ì	WNRTEPLEIMPHIYNHLIFDKPD
						KNKQWGKDSLFNKWCWENWL
						AICRKLKLDPFLTSYTKINSRWI
6838	37206	1	6888	1	1677	THE COLUMN TO TH
0030	3/200	Α	0000	1	1077	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	1	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
6839	37207	A	6889	2	2400	SMRQKENKDIQDLNSALHQAD
						LIDIYRTLHPKSTEYTFFSAPHCI
						/RTYSKIDHTLGSKALLSKCKIT
	Ì					EIITDSQTTVLSELRIKKLTQ\NR
	-					SATWKLN/MLLNDYWVHNEM
						KAEIKMFFETNENKDTTYQNL
						WDTFKAVCRGKFIALNAHKRK
						QERSKIDTLMSQLKELEKQEQT
						HSKVSRRPIPSSEIEAIINSLPTK
						KSPGPDGFTAEFYQRYKEELVP
						FLLKLFQSIEKEGILPNSFYEASII
i						LIPKPGRDTTEKENFRPISLMNI
						DAKILNKILANRIQQHIKKLLHH
		1				DQVGFIPGMQGWFNICKSINVI
						QHINRTSDKNHTIISIDAEKAFN
						KIQQLFMLKTLNKLGTNGMYL
						KIVRAIYDKPTANIILNGQKLEA
						FPLKTGTRQGCPLSPLLFNIVLE
					İ	VLARAIRQEKEIQGIQLGKEEV
						KLSLFADDMIVYLENPIVSAQN
						LLKLISNFSKVSGYKINLQKSQA
}					:	FLYTNNRQTESQIMSELPFTTAS
						KRIKYLGIQLTRDVKDLFKENY
						KPLLNEIKEDTNKWRNIPCSWV
						GRINIVKMAILPKVTYRFNAIPI
						KLPMTFFTELEKTILKFRWNQK
		ĺ				RAHIAKTILSQKNKAGGIRLPDF
			1			KLFYKATVTKTARYWYENRDI
						DQWNRTEPLEIMPHIYNHLIFD
						KPDKNKQWGKDSLFNKWCWE
1		l				NWLAICRKLKLDPFLTSYTKIN
	<u></u>	L_				SRWIEDLNVRPKTIETLEENLG
6840	37208	A	6890	1	1515	
6841	37209	Α	6891	1	1428	

T*RMHK\LSSQFSQLEERVSG EDQTNETKDTHRLKIKGWRK QANGKQKKAEAAILVLDKTI KPTKIKGDKQGHYIMVKGSIK GQLTILNIYAPNTGASRLIKQ RDLQRDLDSHTIIMGDFNTRI TLDRSTRQRLNKDIQELNSAI QVDLIDIYRILHTKSTEYTFFS PHHTYSKIDHTVGSKALLSKO RTEIITNRLSDHSAIKLELRIK TQNRSTTWKLNNLLLNDYW WDTFKAVCRGKFIALNIQKR QERSKIDTLTSQFKELEKQEQ HSKASRRQEITKIRAELKERE KSLQKISESRSWFFEKINNIDT LARLIKKKREKNQIDTIKNDK DITTDPTEIQTTIRQYYKHLY. KQENLEEMNKFLDTYTLPRL WEEAESLNRPITGSEFEAIINS IKKSPGPDGFTAKFYQSYKEE VPFLLKLFQSIEKEGILPNSFY ASIILIPKPGRDTTKKDNFRPI MNIDAKILNKILANKIQQHSK LIHHNQVGFIFGMQGWFNTC INIIQHINRTKDKNHMIISIDAI AFDKIQQPSC	SEQ ID NO:	SEQ ID NO: of peptide sequence		SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence		Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
	6842	37210	A	6892	150	1920	NYQCREVLKGPDGAENHGTRT T*RMHK\LSSQFSQLEERVSGM EDQTNETKDTHRLKIKGWRKIY QANGKQKKAEAAILVLDKTDF KPTKIKGDKQGHYIMVKGSIQQ GQLTILNIYAPNTGASRLIKQVL RDLQRDLDSHTIIMGDFNTRLS TLDRSTRQRLNKDIQELNSALH QVDLIDIYRILHTKSTEYTFFSA PHHTYSKIDHTVGSKALLSKGK RTEIITNRLSDHSAIKLELRIKEL TQNRSTTWKLNNLLLNDYWNL WDTFKAVCRGKFIALNIQKRK QERSKIDTLTSQFKELEKQEQT HSKASRRQEITKIRAELKERETQ KSLQKISESRSWFFEKINNIDTP LARLIKKKREKNQIDTIKNDKG DITTDPTEIQTTIRQYYKHLYAN KQENLEEMNKFLDTYTLPRLN WEEAESLNRPITGSEFEAIINSLP IKKSPGPDGFTAKFYQSYKEEL VPFLLKLFQSIEKEGILPNSFYE ASIILIPKPGRDTTKKDNFRPISL MNIDAKILNKILANKIQQHSKK LIHHNQVGFIPGMQGWFNTCKS INIIQHINRTKDKNHMIISIDAEK
[UO+J]J/411 [A U07J]1 1/7/	6843	37211	Α	6893	1	1797	2 22.00

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
6844	37212	Α	6894	2	1880	APNTGAPRFIKQVLSDLQRDLD
						SHTIIMGDFNTPLSTLDRSMRQ
				ŀ		KVNKDIQELNSALHQADLIDIY
						RTLHPKSTEYTFFSATHHTYSKI
						DHIVGSKALLSKCKRTEIITNCL
		ĺ				SDHSAIKLELRIKKLTENRSTTW
		1				KLNNLLLNDYWVHNEMKAEIK
		1				MVFETNDTLPRLNQEEVESLNR
						PKTGSEIEAIINSLPTKKSPGPDG
						FTAEFYQRYKEELVPFLLKLFQ
						SIEKERILPNSFYEASIILIPKPGR
		1				DTTKK/EENFRPISLMNIDAKIL
						NKILANRIQQHIKKLIHHDQLGF
						IPGMQGWFNICKSMNVIQHINS
:						PKDKNHMIJSVDAEKAFDKIQQ
		ĺ				PFMLKTLNKLGIDGTYLKIIRAI
						YDKPTASIILNGQKLEAFTLKTG
						TRQGCPLSPLLFNIVLEVLARAI
		l				RQEKEIKGIQLGKEEVKLSLFA
						DDMIVYLENPIISAQNLLKLIGN
				1		FSKVSGYTINVQKSQAFLYTNN
						RQTESQIMSELPFTIASKRIKYL
						GIQLTRDVKDFFKENYKPLLNE
						IKEDKNKWKNIPCSWVGRINIV
						KMAIPPKVIYRFNAIPIKLPMTF
					!	FGMIFLSQKYGHVTLLFRLLQW
		l		1		LLNVARNNNKVFYKIYKVGHK
						LSSAYVLCLTCSHFSVSFL
6845	37213	В	6895	13	1506	
6846	37214	Α	6896	1	6635	MTPESRDTTDLSPGGTQEMEGI
						VIVKVEEEDEEDHFQKERNKVE
						SSPQVLSRSTTMNERALLSSYL
		l				VAYRVAKEKMAHTAAEKIILP
						ACMDMVRTIFDDKSADKLRTIP
						LSDNTISRRICTIAKHLEAMLIT
		l				RLQSGIDFAIQLDESTDIASCPTL
						LVYVRYIPLKLRLGGLLQPEKPI
						VLKVESRDGTIQYNSQQHINLS
						AAWLKTAVQGREAPGKQPSKQ
						QHSAKRKTHRTQLKKESGDGP
						HPKFGGGNLASP
6847	37215	В	6897	1	3045	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide		Amino acid sequence (X=Unknown,
NO:	of peptide		in USSN	location of first		*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide sequence	of peptide sequence	deletion, \=possible nucleotide insertion)
6848	37216	A	6898	11	2764	MARELHDKCTSFSSRFDQLEES
0010	3,210	1	0050	1	2701	LSAMEDQMNEMMREEKFREK
		1				RRKRNEQSLQEIWDYVKRPNL
						HLIGVPESDRENGPKLENTLQDI
						IQENFPNLARQGNIQIQEIQRMP
		1				QRYSSRRATPRHIJVRFTKVEM
						KEKMLRAAREKEIQTTIREYYK
			1			HFYANELENLEEMDKFFETYTL
		1				PRLNQEEVESLNRPITGSAIEAII
		1				NSLPTKKSPGPDGFTAEFYQRY
		1				KEELVPFLLKLFQSIEKEGILPNS
	Ì					FHEASIILIP
6849	37217	В	6899	1	3549	FREASILIF
6850	37217	A	6900	387	541	
6851	37218	A	6901	1	545	
6852	37219	A	6902	71	370	RRCKVRPSARGVLRYAVCVCV
0832	37220		0902	1	370	CVCVYPCVHVCTCVRMCLCVC
						VCVCE/CSVEIHCGGQGCARCIC
						LQPQEGGCTEQAVALHCAAQL
						WRNRLKLMKQPVARV
6853	37221	A	6904	1	738	With the first of the transfer
6854	37222	A	6905	11	567	
6855	37223	A	6906	li -	1416	
6856	37224	A	6907	3	1267	
6857	37225	Α	6908	1	552	NALPGRKSRVFSGCVSADTWK
						PPSSAKVTETKGASPAFLRAGQ
		1				PRLVPGETLEKSLGPGKDPQVE
						PQHPVHLPGISSEGFPWDGFNE
						QTPKD\LPNRDGGAWVLGYRA
		1				GPACPFLLHEEREKSNRSELYL
	}					DLHPDHSLTEQDDRTPGRLQA
					<u> </u>	VWPPPKTKDTEEKVGLKYTEA
		1	ŀ			EYQAAILHLKRE
6858	37226	Α	6909	1	561	
6859	37227	A	6910	219	1343	RRLCHTQPTLGMRERSELVNV
		ŀ				CVHSVFSLPSVQAMQEKDEAK
						AETIQALYHQTLEALQTLLKAL
						FIEDPTPAGLKSILEALGPWMNS
		1				GKAHERARAVNTNVSVLNHM
						LLTLPFFMPLGFPALGLLLGRLI
						LHIGDPDEEIGCEALDGIIILYTI
						LELQKRARDKEETNKKELYES
	1					NKHFLGPYNPVSPCQNILRVIEE
						FGDFLGPQQIKDLLLAALEGLK
						GSSEAPGKDSREMMQLASEVM
						LSSVLEWYRHRALEVIPEIMQGI
						YMQLSHIQEPRARQVALLPVSL
						LASSFMTEVVVALLMCPLPLNR
						QPAACASSCPSTAAWAV*GAST
		1	1	1	I	L CCCUMCCERCITTE A CTCL A A W
ŀ	l .					LSCSWPCSFRSITTSASTCLAAW

SEQ ID NO:	SEQ ID NO: of peptide sequence	Met hod	SEQ ID NO: in USSN 09/540,217	Nucleotide location of first codon for peptide sequence	Nucleotide location of last codon for last amino acid of peptide sequence	Amino acid sequence (X=Unknown, *=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
6860	37228	Α	6911	3	730	
6861	37229	A	6912	2	1218	ERKCMQGKYAGAMESEPCVCT EADFDCDYGYERHSNGQCLPA FWFNPSSLSKDCSLGQSYLNST GYRKVVSNNCTDGVREQYTAK PQKCPGKAPRGLR\VVTADGKL TA\EQGHNVTLMVQLEEGDVQ RTLIQVDFGDGIAVSYVNLSSM EDGIKHVYQNVGIFRVTVQALV RLPPHISQCDEVFRFFEARPEDV NPPKDQALLLGELGPAVELCLK EERFADAIILAQAGGTDLLKQT QERYLAKKKTKISSLLACVVQK NWKDVVCTCSLKNWREALAL LLTYSGTEKFPELCDMLGTRME QEGSRALTSEARLCYVCSGSVE RLVECWAKCHQALSPMALQD MHVISTDENQVFAAVQEWNQN DTYNLYISDTRGVYFTLALENV
						QSSRGPEGNIMIDLYEVTI
6862	37230	Α	6913	1	1659	
6863	37231	A	6914	540	824	KPRLENYVKNAEASGADAINW KKGY/LVMEDEMNEMKREGKF REKRIKRNEQSLQEIWDYVKDQ IYVRLVYLKVTGRMEPSWKTL CRILSRRTSPI
6864	37232	A	6915	1	939	Ortizoration I
6865	37233	A	6916	285	458	QVFSQLEGRIALGKFQRPSSLPT GKWTQRCWWGAQWE*D*LFG LHGWWVRPVIAGFP
6866	37234	Α	6917	32	171	
6867	37235	Α	6918	294	461	
6868	37236	A	6919	2	255	LIPCPATVNNVSDIMGMYMKE CSASLITREMQIRTTIMRHHLTS IGTEMIRK/SKNNECWSGRECQ TVGAGQCVRAPCASRSRARH
6869	37237	Α	6920	2	393	
6870	37238	В	6921	87	633	
6871	37239	A	6922	78	260	TQYLRWKCRNHRLLRRSCWEL *TRAVPIRPSWLQNRYQGTSRF CVCFEESSGTPSTQLEI
6872	37240	A	6923	1	369	FVLHEKCLGPFIWTGNTNVEM VKKICAPT*TKHVVW*LES*M WEEPRWPN\GKAPVYSSQREQR RRRVISAFPSEGVLPSTFKNKKF TVKQPQVGRSVGIPEEGIVIIGD GQLQGCLLPLKTF
6873	37241	A	6924	3	201	LKEMQINTTMRYHS\TPSRTS*\I KKTDNNKFWMRMTEKLETSY CWWECETMQPRWKVSLQYLL PLI

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence	hod	in USSN 09/540,217	location of first codon for peptide sequence	codon for last amino acid of peptide sequence	*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
6074	1272.42		16005		12064	
6874	37242	A	6925	1	2064	
6875	37243	Α	6926	95	752	DALLON AND CONTRACTOR OF THE PARTY OF THE PA
6876	37244	Α	6927	2237	5072	RAKSPANIIMTGSNSHITILTLN
					1	VNGLNSPIKRHRLASWIKSQDP
						SVCCIQETHLMCRDTHRLKIKG
						WRKIYQANGKQKKAGVAILVS
	ĺ					DKTDFKPTKIKRDKEGHYMMV
						KGSIQQEELTILNMYAPNTGAP
						RFIKQVLSDLQRDLDSHTLIMG
						DFNTPLSTLDRSTRQKVNKDTQ
						ELNSALHQADLIDIYRTLHPKST EYTFFSAPHHSYSKIDHILGSEA
		1				LLSKCKRTEIITNYLSDHSAIKL
		1				ELRIKNLTQSR
6877	37245	A	6928	3	331	ELKIKNLIQSK
6878	37246	A	6929	364	853	
6879	37247	A	6930	2	531	RPRVRELVGGSAIQALKEWNTT
						GKLRINH*KRCGSTQHHGSKYA
		1				TDKEERPD\ACPLQPQPPQIQNG
		1				PMNGCEKDSSSTDSANEKPALI
						PREKKISILEEPSKALRGVTGPNI
		1				EKSAKDLQRCTVSLTRYRVMIK
						EEVDSSVKKITAAFAELHNCIID
						KEASLMAEMGTVNEQAMRC
6880	37248	Α	6931	1	456	ISILEEPSKALRG\VTEGNRLLQ
						QKLSLDGNPKPIHGTTERSDGL
						QWSAEQPCNPSKPKAKTSPVKS
		ŀ				NTPAAHLEIKPDELAKKRGPNI
		ŀ				EKSVKDLQRCTVSLTRYRVMIK
						GEVDSSVKNIKAACAEFPICIID
6881	37249	A	6932	333	964	KEVSLMAEM\DKVKGRKPW
6882	37250	A	6933	240	691	
6883	37251	A	6934	1	2076	
6884	37252	A	6935	14	485	RRSLESVLSRKLNPF*KATGTSR
						SESAVWAIIDAVSSPQKRLLDSE
						FIDPLMNKKARISHLTNRVPPTL
						NGHLNPTSEKSAAGLPLPPAAA
					†	AIPTPPPLPSTYLPISHPPQIADS
			:			HSPAAPVQGIQPFLTADQYWLE
						NRYPSQHLSRVTASRARHQT
6885	37253	Α	6936	1	643	
6886	37254	В	6937	1	408	
6887	37255	Α	6938	2	197	
6888	37256	A	6939	103	298	CLWLFQEEEEEEEEEED*EEE
						EEE/EEEEEEEEEEEEEE
						EEEEEEEEEEEEEEKI
(000	27257	_	(040	117	267	FLGHRVGI
6889	37257	A	6940	1117	257	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide	hod	in USSN	location of first	codon for last amino acid	*=Stop codon, /=possible nucleotide
	sequence		09/540,217	codon for peptide	of peptide sequence	deletion, \=possible nucleotide insertion)
				sequence		
6890	37258	Α	6941	46	219	LKECNCIVCNSKNKCLKG*KKK
		1		İ		EEEEEEEEEEEEEEEEEE
						EEEEEEEEEEGRRGRGRGRR
						RRRRRRKKK
6891	37259	Α	6942	1	1314	MVSRIWSWISEAQCPLCRPSGT
	1					YGPGEAALLLWLCIEAAAAAA
						AARSTTTPIITNHQSNTIAEQNP
						PRHENGDPKTYIIHVLAEQVLE
			Į			RLPHGVALRHDPLAPVVSRAG
		ŀ	Į.			GVGHQRCAADDALQPLLQGRP
						EPGLAERHGVQDNLILKVSHTH
		1				TQKSAEKRAVSDPSSTVHRPQP
		1				LPPSTVHQSTIPSNPDSSSAYCL
		1	İ			PSTRHGFSSYTDSFVPPSGPSNP
		1				MNPTIGNGLSPQRVHYQHLPTH
		ļ				AGGKRNAALEYGVPRKYYHM
		1				LKITASDYEKTKPFQVGDKVM
						GSGWRKVGMQT*FL*LKKEEE
						EEREEEE\EEEEEEEEEEEE
						EEEEEEEEEEEEEEEEE
		1				EEEEENSLSLTE
6892	37260	A	6943	1	160	
6893	37261	Α	6944	1	306	
6894	37262	Α	6945	1	933	MWLARELEESLAVSLQGEPGL
		1				LAGAGRLQGSPFSHPMEPGRQS
						ETLSQKKKKKEEEEEEEEEE
		1		i		EEEEEEEEEEEEEE
						EEEEEEEEEEEGEEEG
		1		•		DQLDTM\FGTHQQTSPTQRSAR
			İ			RRPCSVQGPRS*SPFSHPMEPGR
		1				QSETLSQKKKKKEEEEEEEEE
						EEEEEEEEEEEEEEE
						EEEEEEEEEEEEGEEEGEE
		1				EEDQLDTMLWDSSTNLTNTAL
		1				SKEKTMFSSRAKIVKPNGEKPD
		1	ļ			EFESGISQALLELEMNLDLKAQ
			}			LWELNITAAKEIEVGGGRKAIII
		1	ļ			FVPVPQLKSFQKTQVQLRRILP
		1				KPTQKSCTNNKQKLPRSCTLTA
		1				VHDAILEDLVFPSEIVGKRIHVK
		1				LDGSHLIKIHLDEAQQNNVEHK
						VEPFSGVYKKLMGKDVNFEFP
6895	37263	A	6946	1	1365	
6896	37264	A	6947.	416	610	EFKKPVLEGGA*RGEGGSPCV
						WSFFDPVHFNYQAGLFRVALP
						GEQCRTGRLVARVYTGISGASF
6897	37265	A	6948	1	846	

SEQ ID	SEQ ID NO:	Met	SEQ ID NO:	Nucleotide	Nucleotide location of last	Amino acid sequence (X=Unknown,
NO:	of peptide sequence		in USSN 09/540,217	location of first codon for peptide		*=Stop codon, /=possible nucleotide deletion, \=possible nucleotide insertion)
			, 	sequence		, , , , , , , , , , , , , , , , , , ,
6898	37266	A	6949	3	620	HVTLLIVVTQSHLVIHNLFEGMI
		l				WFIVPISCVICNDIMAYMFGFFF
		1				GRTPLIKLSPKKTWEGFIGGFFA
						TVGFGLLLSYVMSGYRCFVCP
						VEYNNDTNSFTVDCEPSDLFRL
						QEYNIPGVIQSA\IGW\KTVRMY
						P\FQIHSIA\LSTFASLIGPFGGFF
						ASGFKRAFKIKDFANTIPGHGGI
		İ				MDRFDCQYLMATFVNVYIACV
						LRPL
6899	37267	A	6950	50	554	
6900	37268	В	6951	446	625	
6901	37269	A	6952	1	2149	
6902	37270	Α	6953	1	1704	# 1 / Will /
6903	37271	Α	6954	1	1853	
6904	37272	A	6955	1	1305	MVKGSIPQEELTILNIYALNTGA
						PRFIKQVLRDLQRDLDSHTIIMG
						DFNTPLSTSDRSTRQKVNKDIQ
						ELNSALHQADIIDIYRTLHPKST
						EYTFFSAPHHTYSKIDHIVGSKA
						LLSKCQRIEIITNCLSDYSAIKLE
						LRIQKLTQNCSTIWKLNNLLLN
						DYWVHKEMKAEIKMFFETNEN
						KDTTYHHLWDTFKAVCRGKFI
						ALNAHKRKQERFEMDTLTSQL
						KELKKQEQTHSKASRRQEITKI
						RAERKEIETQKTLQKINESGSW
						FVEKINKIDRPLARLIKKKREKN QIDAIKNDKGVITTDPTEIQTTIR
						EYYKHLYANKLENLEEMDKFL
						DTYNLRRLNQGEVESLNRPITG
						SEIEAHNSLP/TKKSPGPNGFTA
						KF*QEELTILNIYALNTGAPRFI
						KQVLRDLQRDLDSHTIIMGDFN
		l		}		TPLSTSDRSTRQKVNKDIQELN
						SALHQADIIDIYRTLHPKSTEYT
		ŀ				FFSAPHHTYSKIDHIVGSKALLS
		l				KCQRIEIITNCLSDYSAIKLELRI
						QKLTQNCSTIWKLNNLLLNDY
						WVHKEMKAEIKMFFETNENKD
						TTYHHLWDTFKAVCRGKFIAL
	1					NAHKRKQERFEMDTLTSQLKE
						LKKQEQTHSKASRRQEITKIRA
						ERKEIETQKTLQKINESGSWFV
						EKINKIDRPLARLIKKKREKNQI
						DAIKNDKGVITTDPTEIQTTIRE
						YYKHLYANKLENLEEMDKFLD
l	ı		1	1	i	